

<110> Byrum, Joseph R.

<120> NUCLEIC ACID MOLECULES AND OTHER MOLECULES ASSOCIATED WITH PLANTS

<130> 38-21(15598)B

<160> 36935

<210> 1

<211> 147

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 1

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atggatggtg gnttggaca tttggat 147

<210> 2

<211> 378

<212> DNA

<213> Glycine max

<400> 2

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agcaaccgtg aactgctccc tctgcacttg ggttttaaa tcccactgaa actgacaacg 180
ccaacctcac tggcgctcc ctctcaccca aatcgccacc tcgagtggcg cttccagct 240
aggaccctgc aactgcaggt gctgcaacct atggttgcag aagtgcaggt gctgaagtgc 300
ctggtcaaac cagcgcttc atgcacccctt ggacgcgccataagagctg cgatatgcag 360
tcctttgtcg ccagtcac 378

<210> 3

<211> 423

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 3

agcttgtgag ttgtgagtgg tactactgaa aataccccc ccataccccc tttcccccta 60
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tctaggtaa ttccatcatc taattacctt gcagacccta aaatcagaga agattgagtt 180
tgttgtagct ccatatgaag cagatgctca gttagcgtac atgtctcagc ttggagtaca 240
aaatggcgg gttgcagcgg tgatcacaga agatagtgtat ctaatagcat atggctgtcc 300
agctgttaa actcctccaa tactgtgata ttgcgcattgg aggtttactg cnntttgat 360
atctcgattt atttacttgt tcactattca gcttcataaga aagcatgcat tttgggatata 420
aat 423

<210> 4
<211> 462
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 4

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gcattttcag ccaacccaag gcaccacttc ttgcaacaca ctactggcct acacggatcg 120
aaacagtaca gcagatgaaa atggtgggtc gcattaacct tcacaatttgc gcagcacaga 180
aatcgacta ttttctcaa cgaaccattt aatggaagca gaatgaatga ggatgcagtg 240
ttactggctt ggtcatggtt cacagagctg gagaaaagat ttacagagca ttacaactac 300
tggccagca acctatcagt agctntctgt aactagcata gatggaaagc tattgaccaa 360
atgtaacaat gtatctagtc tggattctaa cagaggacc ttcattccac acagtcatac 420
tcctgtattc ttagtacccc ggtactttc tataatataa at 462

<210> 5
<211> 394
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 5

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tcaaattccta aaatttgagc tccttagggag caaaacaatg tgtgttcctt agagaggca 180
tcagctacca catttggttt tcccttttg tatttgataa catatggaaa ttgctctagg 240
tactctaccc atttgcattg cctcttggtt aacttgcttt gccctctaattt gtacttaagt 300
gattgatgat cactatgaat gacaaattcc ttggaaacaa ggtgtcgaa cctacccttc 360
ngcgggaggg cgacgcgtga ctgcgggat gcgt 394

<210> 6
<211> 464
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 6

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agttacaaca agtgttacac atgcttctat ttatagacta ggtagcttcc ttgagaagct 180
ttcttgagga aacttccttg agaaacttct ttgaaaaaac ttcccttgaga aggttagagct 240
tagctacaca cacccatctc ataactaagc tcacccctt gagaagttc cataagaaga 300
ttcctaaaga agctagagct tagctacaca tacctctcta atagctaagc tcacccctt 360
gagatggaa gctagagctn tgctcacacac ccnctatgtat agctaagctc acccccattga 420
caaaatacat ganaatacaa aaaagatccc tactacaaag acta 464

<210> 7
<211> 373
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 7

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tccgcttaagc gcaacactca ttggcttaagc gcaaggaaga atctggaaaga aaatgagctg 120
tacaagttcg cttagcacac tgtttcgtct cactaagcgc accgcttcag tccatcagct 180
aagcgagaaa ggcacgcgt aagccgaaat tcactaatgt gcgcttaagcg gtccagaatt 240
gcgcttaagtgc acgagcacg aacaaggcca cctatttaag cttgaaatca gattttgtga 300

agggagtttgcgcttagattcagagcttgcatgtctaga gattctagag agagaaagg 360

ccaatccag aga 373

<210> 8

<211> 462

<212> DNA

<213> Glycine max

<400> 8

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tacctgtcgcaagggtttgtgtttgtctctgtgtccaccatacagacccccc 120

cttccatgca gcaacaccttggagcaattgagc agcctgaagcttatgtgcataatattaca 180

atagacaccttc tcaacaccttgcagcagcaaaatc aaccacagtagtgcgatattatgacccccc 240

gcaacagata caacccttggatggagaaatc accctaacctt cagatggtcc agccctcagc 300

aacaacaaca gcagcctgtcttcccttccaaaatgtgc tggcccaagc agaccataca 360

ttcctccacc aatccaacaa cagcaacaac cccagaaaca gccaacagctt gaggccccctc 420

cacaacaccttc cctcgaagaa cttgtgaggc aaatgactat gc 462

<210> 9

<211> 421

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 9

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taatcaataa tctatctttt acataatctt cttcaacac cttcaatca atctttcaat 120

atcttcttta atctctttca acatttcaa cagatcttc tgatttattt cccttcatct 180

ttctaaaagt ttttgttcaa tagttctct tccaagaaaa gttcttgcgtt caaaaacttc 240

agctattcat ctttttcatt ctcttccttcc ttgcacaaaa gaaggaagga ctaaccgcct 300

gaatttttt gtgtctctct tctcccttac aaaagattca naggactaac cgccgtat 360

atcttttgcgtt tccccatata aagattaaa ggactaactg cctgagaatt cttgtccca 420

a

421

<210> 10
<211> 404
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 10

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cagatgcacc accaccgcct ctacatcagc ctccgtcctl agagtccatc ttagctcaca 120
tgcagaggat ttagctccat atgcattcat atatgcaca tgtggccgac caataggcg 180
ccaatcatag gggataggtg cagctgaatc agagctttt ccagtacacc ctacgtcagc 240
agagcttagga tcccagccct tactcgtggc ttactcccga gtagtttggg gccacagttg 300
catggcctgg agataggccc aattttcaag tagggacaag accctcaaag gccccaggag 360
ttgaagatgg agctcaagaa gacgacgaca tangcgatgt gatg 404

<210> 11
<211> 284
<212> DNA
<213> Glycine max

<400> 11

agctttccaa gatattaagt ttttcctcag aactgtcgta agcgaagatg ccaatgtgct 60
atataacaact ttctgttgcc catctgcttg tggagacat gtggctgaaa ataacaattt 120
agtgcaccaac ttgacccaca caggactacg caaatggctt atgaacttac agtccttatac 180
actaacaatg ctgcttgta aaccatggat gctcacaatc tccttgagga acaaatttagc 240
cacatggaa gcatcatcta cttcttaca tgaaataaaa tgag 284

<210> 12
<211> 373
<212> DNA
<213> Glycine max

<400> 12

tcgatgaaga tgaaccatct aacaatgaac aagggtgtat tcaccatttg gtgtgttagct 60
actaatattt taagtcctat atataaatct tctctttgag cacttcttta tagctaatgg 120
gaagtacatt ataaccacca atttgataag atattgcgac ataggcaaac cactaatcca 180

atgtacatac tgca~~gt~~cat acatgtt~~gt~~a tccggaaagg attcacatgc atagagacat 240
tgtgaaccca agattcctac tatgtt~~gt~~t gcaatggaaa gagttaacaa acagtgtt~~ga~~ 300
aaccacccct tgttaatgcc tatgaagaca aacttactgt cacacccata ctaaaacacc 360
cctaacat~~at~~ act 373

<210> 13
<211> 339
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 13

agcttgcatc actttac~~t~~ tttcattcc atcgta~~c~~tct tgcttcaaaa actgccc~~a~~at 60
cttgggtg ttttagt~~gt~~t cattattcta atcaacatag tggatgat~~at~~ tgctccaaac 120
aagattgctc ttgc~~ctt~~ga ct~~t~~tcttat ctcc~~t~~tcgn gat~~tttttt~~ attt~~gag~~caa 180
ccg~~tt~~gatta tccggtaggg gtggaa~~ct~~tc gtat~~at~~gtct ttaat~~at~~t~~tt~~ cccat~~ag~~atc 240
acaagcatca agatagg~~gt~~tt ccgtt~~ct~~aat agc~~ct~~agg~~tt~~ tggtaat~~gt~~t ntcc~~at~~tgaa 300
tagt~~gaa~~agc ctat~~gaa~~gca cg~~gac~~accc~~c~~ct agtcc~~c~~tta 339

<210> 14
<211> 397
<212> DNA
<213> Glycine max

<400> 14

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gataattcga gat~~cac~~t~~tg~~a aatt~~at~~gaa aaaaatt~~gtt~~ tccgt~~g~~aaga aaatt~~ca~~agc 120
caaggcgtt ccgtgggt~~g~~a tttcgc~~g~~aag atttcaacc gttcttcgac gttcttcg~~t~~tt 180
cg~~t~~tcttcgt cgttcttcag tcttcaaccg gtaagttccc gaaat~~cg~~aac ttttcaattc 240
attctat~~gt~~a cccttagt~~gg~~ tcctcattt~~g~~ tttcac~~g~~cg~~c~~ cttttat~~ttt~~ cg~~ttt~~cattt 300
actttccgta cccc~~ttt~~g acgt~~g~~c~~t~~ta gtcatttact taagt~~cat~~gt tctcgcc~~t~~ta 360
tcaaaaaata aaataaaat~~at~~ ccact~~gat~~ca ttt~~gag~~t 397

<210> 15
<211> 384

<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 15

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gagttgggaa ggattgatn tggaacttgt cgtggtcag aagtttagttc aagtgcgaac 120
actactagaa aaagagctt ttgcgtatca ctacgacat cggccaaca aaactgtcga 180
agtatattaa atggtgcatt tgtgtatca caacgaaatgt gtgcacccat ccaattttat 240
ggttgacatt ggcacaactc ctttgaagg ttttggagg gactcgagag tgaggaaact 300
agacagctt ggttattctc gattccgtt acatacttaa tgctctcaca acatagtgaa 360
gttagggtag taaaatttac catt 384

<210> 16
<211> 243
<212> DNA
<213> Glycine max

<400> 16

cgaaccgcac cctactttat acggcgacaa acatgtggat atagacaaac atgcgctgac 60
ccgtctcagt gtcatgccta aggctagctc agcatgagtc caactttatc tagcgcgatt 120
cataatgagt tgtgccacat tttgcctata agtaggttag gcgattttt tcaaccaatt 180
agactctaattt ccatggtgaa tcaagtttac tcacaataat aataataatc ttttactta 240
cct 243

<210> 17
<211> 331
<212> DNA
<213> Glycine max

<400> 17

agctatatat cttttcttct tggttctgt tgctctgtat tgctttgggt ctatgtatc 60
cttttatatt tcatactatc tttgacacat gggactaaac attgaacagg tggaaagggt 120
gtccagccaa aattccatgg tcctttaaag aagaagagac tatgtatctg ctggaggaa 180
aagtgagggt tactgttcaa gggctgttg ggtctttga aattgggggt ggtgatttag 240

ttgtcttccc aaaaggaatg aacattactt gggaaagtatgat tgaaactgtg aagaagcact 300
acagcttgaa aaaataatga tgtgtactta t 331

<210> 18
<211> 443
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 18

ctnggtgttg ttccatttgt gcgagttact gaggtgcaat ttcaatttttta attggataat 60
gagaaatgtt agcaatatac taccgtatga cactgcatca cacactttat tatttgccac 120
aatttattgg aaatcacaaa attttgcggg ttctgttact tatttaatga acttcactcg 180
tgattttgga atttctaata aattttaacc aataataata ataataagat gtgttactta 240
gaagggcatt gtattgctag cactccttgc gaagtatagc atacaacat gaaaggaatt 300
ccattttaag tattatcctg taccanaacc tcacttttagt ccccaattttt ggaaatcaca 360
gttctttca ctgacaaaatg acttacagtt nttagttaaaa atagggatta acaagagtgg 420
agcataacaag accaggaggg act 443

<210> 19
<211> 396
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 19

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tcttagtatt tctttcaagt attgtgccta ctctagaaac aagaataaga gaaaattaaa 120
ggaaatgcag ttagtcacat gatatattga gatttgcggg tttttcttt ttttgcattt 180
gataaaaaag acaaaattgg tgcgtttgc attaaggggc ctttcagaag aacttgcgaa 240
aataaattga tcagttaat ttccattatac ttcaagtgaa aagattttta tactatgaac 300
taaccaaaaa tcatcctatg attnttaata taattattat aaaattacca tacatcataa 360
tttgcggatg tagaanacat aaacaacgtt tacact 396

<210> 20

<211> 459
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 20

gtcctcgccc cattcctgcg aaggaaaaca tttggatagt tagtntacc aagaaatgct 60
acccttaaaa caaaaatggc atacaacctc ctccaataaa tacaaacatc aatgtaaatt 120
tagagcaagc ttatgcgcac attttcttac gaacattcac tcgcacaaga tattcttcta 180
actaagaaaa atgcacccat gcacaatcaa ggcacttcg ttacctacat tatttgtatg 240
tacttccaag gtgtactacc tacaccacat gcatttcctt ggctaaattt acatacatgc 300
atgctcaaag cctcttgct accaaaagtt gcacacatgc aaactttatg atgaatctt 360
gctatctaca caataaggtg ctacacttca tgcttataat caagtgtttt actaccagaa 420
gccgcacatgac aatgtcagta tattttctt tgccgacta 459

<210> 21
<211> 328
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 21

agtttctccc ccaattttct ataaataggg ggagaagtgt agtagaaaag ggttcagtcc 60
cttaggcact tctctctt tcgaatttgc ttagaaaaat tgttccgtg aagaaaatcc 120
aagccgagggc gcttccgtaa cgtttccgtg agtgattttg cgaagggtttt cgaccgttct 180
tcgacgntct tcattcggttc ttcatcgntc ttcatcgntc aacgggtaag tacctcatac 240
caagcttttc aattcattct atataccgn nnngggccac attatggttc atgtattttt 300
attctcgntt catttactct ttatacc 328

<210> 22
<211> 391
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 22

ggctctanat ntacattgat gtttgttattt atgggaggag gttatatgcc attttgctt 60

taagagtaac gtcccactgg taaaactaac tttccaaatg tttgccttcg caggaatggc 120
cccgaggaag cttgcctcaa agaggccag gaaggacaag gcggccgaag gaactagttc 180
cgccccggag tacgacagtc accgcattttag gagcgttgta caccagcagc gcttcgaagc 240
catcaagggg tggtcgttc tccgggagcg acgcgtccag ctcatggacg acgagtatac 300
tgatttccag gaggaaatag ggccgcggcg gtgggcacca ttggttactc ccatggccaa 360
gtttgatcca gaaatagtcc ttgagttta t 391

<210> 23
<211> 352
<212> DNA
<213> Glycine max

<400> 23

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accttatgaa aaagtatgga agatagtaga agagtgtaga gactcctaga atgtgtggag 120
tattctagag aattagtctc catcctagga tacaagtaat ctccactatt tattgtggag 180
gtggagtagt ataaataaaag gtaggatcct tcattcctaa aaaatctaag tagagagtct 240
ctctgagaga gaagataaaat agctttggaa gtctctatcc tcaaacataa gtaaggctct 300
ctgagagaga agataaaatag cttggaaagt ctctatcctc aagcttgagt ga 352

<210> 24
<211> 438
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 24

ggcacactct ntgattatct tggtctacca agtgtttatt acacaatagt gaaatgcact 60
tattcccatt ctccccgctc aaccactgaa tcgaattatc tccagccacc caactacacc 120
caaaatagag gttcagaaag gaagcaaact aacactgccataa acccccaa gttccaggtt 180
ttaggtggga attatactat tatcaaaacg gttctacaa cctcttatag aagcataact 240
cttgcaaaca ctttaatag taaaaaaagaa aaaaaaaaaact ttacgtcaca gaactcacta 300
ccaagtggaa aaccaccaac attgtcggtt tgtactctgc agtctgcacg tgttccata 360

aacagaaaaca cattgatttt aaattaatta atcgattaat actaccatca agtagtacca 420
ccctatatatt ctttctta 438

<210> 25
<211> 104
<212> DNA
<213> Glycine max

<400> 25

ccacattatt tccatgacac aaattgcaaa atgatgattt ggaaacttca tgcaaaaactg 60
gtcatgcatg cacctatgca gacactcaag tgtcaaattt ttat 104

<210> 26
<211> 386
<212> DNA
<213> Glycine max

<400> 26

tgtctcagcg tttatgcgag acagagacca acatgttagc tatcatcgcc aagtaccaag 60
aagagttagg tctagccacg gcccacgagc atagaatcac ggatgagttat gctcaagtgt 120
atgcggaaaa agaggctaga ggaagggtga tcgactctt acaccaagag gcaaccatgt 180
ggatggatcg gtttgctttt accttgaacg ggagtcaaga acttccccga ttgttagcca 240
aggccaaggc gatggcagac acctactccg ccccccgaaga gattcatggg cttctcggct 300
attgtcagca tatgatagac ttaatggccc acataattag aaatcgtag gaaaattgtt 360
tggtctctca gaccttgact ggatac 386

<210> 27
<211> 379
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 27

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cttaagcact tctctctttt tcgaatttgc ttggaaaaat tgttccgtg aagaaaatcc 120
aagccgagggc gcttccgaaa cgttccgta acgtttccgt gaggaatttc gcgaaggttt 180
cgaccgttct tcgacgttct tcattcgttc ttcatcgttc ttcatcgttc aacgggttaag 240

tacctcgaac caagctttc gattcattct atgtacctgt ggtggccac attgtggttc 300
gtggattttt attctcgntt catttacttt ctatacccc tttgacgtg gcttaagcca 360
tttatttaag tcatttctc 379

<210> 28
<211> 395
<212> DNA
<213> Glycine max

<400> 28

ttggatgcct ataagattat taaggctgaa gtcgagaaac tatgtggata gcaaattaag 60
atcgtgagat ctgattgagg tggagagtac tatggtagat acacggagaa tggacaagca 120
cctagtcgt ttgcgaagtt tgccagaa catggatta ttgcccagta cactatgctt 180
ggttctccgg atcagaatga tgtggcagaa agaagaaacc gaactttaat ggacatgata 240
aaaagtatga gaagtaataa aaaacttcct caattcttgt ggattgaagc attaaagacg 300
gttgtatata tattaaaccg gttccaacc aaggctgtct taaagacacc tttcttagtta 360
ttcaaagggtt ggaaaccgag tttgcgacat atatg 395

<210> 29
<211> 408
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 29

agcttanagg agcactcana tcgggtgtat ttaacccat ggcctagact ccgaagagtc 60
cgtcagggcc tctccctcct gattcaggc caacccanaa aacattnag cacacagact 120
ntatctatga actgtacaaa atacacgact cctcaattgt tctcaaaata attttatcta 180
atcgcgcttg tgattaaact cgtcaggtcc caacagtggt tcccatcata atactcgcca 240
cgcattaaact cgtcgccctt agattcatag ttcacaaatc agggcacaca acatctcaat 300
gcacatatat attacaagtc aatacatact caatttatca catacatttgcgtcaatca 360
cagtggata atctcaattt aacatgttat cacacctcat gaatcata 408

<210> 30

<211> 443
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 30

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agtctgatat ctcaagatta catttgaata aaatcatgag gttgaattaa ttgaaatact 120
ttaaggat agcaaaaaag gtttcagcta aaacaaatgc aaggcagcgt aagaaataaa 180
ttactacatt agcaataacg cttaaatatc tacataaaca gaatattcca taaagattat 240
attnaagccc catgctgaga tgcaagtaat atgctgttc atatttatca aaatataffa 300
atggaaatga tgcaggaggc ccacagaatt aagtcataaa cctgaactca actacatctg 360
tgcatcacaca taaaaccanat cctaccatn taattntaca ccctccccan acccacaatg 420
aatatggcct aaggaataca tcg 443

<210> 31
<211> 398
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 31

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tattcattag acatagcgtc atctatgtac tttggcttga tctcgaaaag caaagttcta 120
tgcttggaaag agtccttgt caggactttg tccttaggat caccaatgtat ctgggactct 180
agatgatctt tttgttagcaa tcatccagtt ggttctctga cttatagagg ttgatcatcc 240
actggtgagt tggacgcata ttgggtcttga ctggacctag caacatattt cacgatattn 300
tctactttca tctcagtaga agactcatct agctccaaca ttgttagtggt aggcttattg 360
tcattaagtc ttacgtgaat ggcctttcc acagtc 398

<210> 32
<211> 458
<212> DNA
<213> Glycine max
<400> 32

tcaccttgc cgcctcctca tagttgtgc atgagacaac atgctctatt ttcatctccc 60
actccaagta ggcctccgga tcattttc cttaaaggg aggaatgtt agtttaatac 120
catcaattcg gcttgacta ggaacaccat cattccctct tctcctcctt tcttcttcat 180
tatgatctct atttccatt tgatccaacc tctcatggag cacatcatct cattgttca 240
ttaacctctc caaatgatgc atcacagctt gcatttggaa ttgcgaaagc cccactccat 300
cattaggatt tggcctgcc atctcataca aacacatcag acgtatcaag acaattata 360
ttgctgttg aatacctcac tcactcaagt gtatcacaca attatggtt ttctctaattg 420
aacactctt gccttctacc actctaattc cacttgag 458

<210> 33
<211> 382
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 33

agcttgata taatgcttcc ttgttattta taggtataga ggatcttagcc acttgagagt 60
gtcttgc(cc) gactcagttac aatgcattat attagcgcga catagcttg gtctacattt 120
aattagtata tatagttaca ctttattcaa tccaagaatg gaaaaaggaa agcagtgg 180
aagaatgatc atgggtcaag cttgagtcaa gtaatctgat accaaaccga atgaaaatgg 240
ttaggttag tttttttttt gtgactgtct caaccacaac taatcacaag tggatgtgc 300
ctaagtacg tcattatatt aaagattctg tgaatgaagg aaaaaaaac acacanaaat 360
agggtagag tagggaaaag gt 382

<210> 34
<211> 460
<212> DNA
<213> Glycine max

<400> 34

cctgagtgaa acaatgagac tcttcacagt taaatttggaa ttgcacgtt catggacact 60
ggtaatcgat taccaaaaca ttggatcga ttatgcctt ttgaatatat tggaaacgtt 120
gtaaattcag tttggaaaact ttttcaact cttttagcta ctgagaatcg attacaacaa 180
tatgaggatc gattaccaga gagaaaaagc tctttggtaa agatggtc aaaaactcac 240

gagctataca acgttgagaa aaaacccccc taataacttat attgatagag tgtttataac 300
attctcaaata gttgaatgtt gaatcttgat ctgttgcgtt gagaactcga gtattgagtc 360
ttgatttata accttgatgc ttgtatgtt acatcatgaa tcgtaatct tgataacttat 420
ctgaaggctt tcttcttgag tcttgaattc ttgattcttg 460

<210> 35
<211> 320
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 35

agctttaga gcttgagttc taagaatgag ctgagttatt ttgaaccat tntgctgttt 60
atttttccta aaatggatta tgaatatagc ttggatacta tggaaacaaa aaatgagtat 120
tctgatgaag gtgaagtagt accgggtta tccattaatc cggttagttt ttgcgcAAC 180
attgttccaa tgattaatgg taaatatggt ttgcattacc aaagtgaatg caattctaca 240
aagaattgca gtcctattgg gggatgccca accaattgcc ttatattgtg tctttgaaag 300
aactggtttg ttagatgtaa 320

<210> 36
<211> 368
<212> DNA
<213> Glycine max
<400> 36

tcacattcac tatcctctac atcatattca aacttgacca aataaatagt acagtcatct 60
cgactcaaag aaggacatct aagtctcata caattaatat agaacctata tcctaattgtc 120
acatcctatc aaagcgtggc gctaccgcgt cctctagctt gaggctcttc atagtcatcc 180
acctattcat ctgctacccc gaacacagag cttgagatca tcacaggatg cgaacacaaa 240
cagcacaccc ggagtgagat atcacacttt taactactat agagaaacaa cacaacatct 300
atgagccgaa gacgatttac ttaccatata tcacattatt tcatgacttt gtccttcatac 360
gatcacac 368

<210> 37

<211> 414
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 37

agcttatcat ctatTTatac tacacatAGC aatgaataCT tattCTGGAT cactGCTTAA 60
tGaaATTaaG gatCTGTtta ttTACATTG taaACAGGCA aatgtattAA ttTAAGATT 120
gtatCTAAAT tGttCTcaAC tatATAAAT tataACATT ngtattaACA tGtATAAttC 180
tagcatacAG gcAGTTtta gaaATATAcc atgtGACTCG atCATGTGTA tGTTTGAGTT 240
gataATCTC ttggaggAGt gttaAGATTc aacACATTG tAtTAATATG atgtGAGAGt 300
cacAAATTNT tATCATTTC aATATAATTc aatcaATAAC aaAGAAATAA ttTAAGAGA 360
ctcaacAAAT atgtCTCTAC tGTTCTCAC ATTATAATA taatcaATTG CTG 414

<210> 38
<211> 288
<212> DNA
<213> Glycine max

<400> 38

gttccAAAGA ggtCTTCGGC attacATTCA aactCGATCC attGTGATA agtACATTG 60
cgaccACGTG tgCCGTACAT atcAccGACG catGTACAGC cttGATGTGC cctCTCCTCT 120
caacGGGAAT aacttCTTGC acgaACGCGA tataATTGCC gatGGCTATA tgATTGGCTA 180
tgCCGATCAG aacGATGCGT gagatATAAT gagCTACACT gtcATGGACA tgATCCATAA 240
tctGAGACGC actGATAcAC tccCTCAATT ctTGTGGATT gaAGCATT 288

<210> 39
<211> 317
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 39

agcttGTCAC tGtGGAATTa accaAGTCTC tgAGGGACGC gttCAAGGAT aattCGAAAT 60
catATGCCAT agcGATCTCT tcACTCTGAC catCAATAAT tactACCATG gtttCTATCT 120
ccaAGATGTC aaccAGAACT gctACCTTT gcCTGAGCTT gaAGGACGTA tatGACACTG 180

aatacaccaa taagtaatta atactacaat ctaatagctt aatactaac agttgacaca 240
aatattaatt ntcattacct taatgttctg aagtgcacgt ctaaagttt gtgcataaca 300
gtatcgccac taaaacc 317

<210> 40
<211> 435
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 40

cactatctat acttagcttgc caatctatgt gcaccagata gctaataagac ttttgttattc 60
ttncgtttag agtatgatat ctcaagatta catttgattc acctcgatg gctgcataaa 120
ttgagttatac ttcatcgacg ttgcaaatac aggttagct taaactgatg cgctgcaccc 180
cagggaaatag agtactacat tagcagtaac gcttaataat ctacattatc agaatattcc 240
ataaaagatta tatttaatcc ccatgctgat atgccagtaa tatgcctgct catattgatc 300
acaatatatg aaggatatg atgcacatg cccacagaat taaatcatag acctgaactc 360
aactacatct gtgcatacac ataaacccaa tcctaccatt gtaattttac accctcccc 420
aacccacaat gaata 435

<210> 41
<211> 285
<212> DNA
<213> Glycine max

<400> 41

agctatatac aagctttct ttcatcagat gctccgaaaa tttaacttct tggtgtgata 60
attagggggg agcagttat aactggattt gtatctgaca gagagaaatc ttaacacaag 120
tcactctgac actcttattt tataacaaat taaggccact gagttgatc cagctatcca 180
aaagctgttag gaataaaaaaa tctattaaga gcaaacacac acctcgaccc gtgttatgca 240
agtaaatggtaatgtt aatgttacgcg actacgcagt agtacgggtt ggttag 285

<210> 42
<211> 452
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 42

ttgatgatag anattaattg aatgatttat tattaattct tcttaatgtt tggacaatat 60
ttccaatatt agaaattaca aactcaaaag aaaccaataa ataaattcct agtaacaaaa 120
cttgggggggataaatttcat atttatcaat atttattata taaataataa taaattataa 180
ttaaaaaatg aataagtatt atgttagata ttttataat aatataagat aatatcta 240
ataaaaaaaaat atctatcaat gagatcggtc acttgtgtta gctaacttac atgaaaagtc 300
aatgagatct gttacttgtg ttgcattgggt gttagacgaaa cttgaacatc attagcaatt 360
atcaagggtc tcctatcatc acataaagta tgggttgat acttaacaat aagcagacca 420
tcacagaaaag gatatgatag cactctgact at 452

<210> 43
<211> 335
<212> DNA
<213> Glycine max

<400> 43

agcttctact tatgtggcag ggcgggcttc ctccacccccc ttgtctccaa cgcgaaacctt 60
gaccattatt ctcccttccc gcgatgcttc ttttcatgtc cgcctgagtg ggcttatagc 120
ctaaaccata ctccccacga ttcccttggg tatttatcag gctagttatg ccgcccgtgt 180
ttttcctaa acccatcccg ggttcataac cgttcccaa cataactcgg gccatcatta 240
tcgctgcac ggacagacaa ggcttccaa agagggagtc cacggaggaa atgctgacca 300
cctcaaaaga ctggaaagca gtttctaacg attct 335

<210> 44
<211> 434
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 44

tgcctcanag agatccaaga aggataaagc agctgaagga actagttccg ctccctgaata 60
tgacagccat cgtttagga gtgctgagca ccagcagcgc ttgcaggcca ttaaggatg 120
gtcatttctc cgggagcgcac gcgtccagat cagggacgac gagtataaccg acttccagga 180

ggagatagtt cgccggcggt gggcatcgct ggttacccc atggccaagt tcgaccaga 240
catagtcctt gagtttatg ccaatgcttg gcctacagtg gagggtgtat gagatatgcg 300
atcctgggtg agggggtag tggatccat tcgatgcgga tgctctcagc cagttcttgg 360
gatatcctt agtgctggag gagggccagg agtgcaagta tggccaaagg aggaaccgg 420
ccgatgggtt tgat 434

<210> 45
<211> 408
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 45

agctntntgg agtagaaaca tgggaccaac tcattttatt tcaaaaagga agtcatatct 60
agtcaaggc tgagagacca tacaagttc ctaacgattt ctaattatgt gggccattaa 120
gtctatcata tgctgacaat agccgagaag cccatgaatc tcttcggtgg cggagtaagt 180
gtctgccatc gcttggcct tggctaaca tcggtgaagt tcttgactcc cattcaaggt 240
aagagcaaac cgatccatcc acatgggtgc ctcttggtgt acagagtcga tcaccctctc 300
tctagccctct tttccgcgt atacttgagc atagtcgccc gcaatcctat gctcgtggc 360
cgaggctaga cctaactctt tcttgcgatg atagctagca tggtggt 408

<210> 46
<211> 74
<212> DNA
<213> Glycine max

<400> 46

tctcaaggaa gtttctcaa gagagcttct caaggaagct acctagtcta taaatagaag 60
catgtgtAAC actt 74

<210> 47
<211> 358
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 47

agcttgaagc atctcatcat aggagatgat gatcacaaag aagcanagaa ggtagtagt 60
aataataata gtaataataa taatagtgtat catgagagag gtggcaaagg agatcaagg 120
aactcacttg gtgaacactt cacagaggaa gagaagcgcgataatctca gctggtagg 180
atgcaacaga ataaggacaa cctccaaggc ttgaagttga agaagtttgt gcgtcggtac 240
gccaaagttt tggngcattt gatgaangct aagcgtgatc ctcatctang tggtgatgct 300
ggaaaaaac ctgtnttcaa gttatcagcc tagccaggaa aaattttgga gttttact 358

<210> 48
<211> 407
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 48

tgcatttcac gagcatcaact ctttntcctt ccctcttgg ttgtgacatt ntcttattga 60
gaatggaaaa ttatccat accttcaaattt tgcaaaagaa atgtatccc aaaatgctac 120
tctaaaatag gaagagagtt tgctcttgg ttttcgctg gtccatataat atactaaatt 180
agaagtcatt aaacaagctt tttcaactct cataattgtg gcattacta ttgaaggtag 240
ggaaatgatc ttaaacggat tgaaaatatt aagcaaggaa aattactgca gaaattctta 300
acaaatgaaa tcagaagtca cttctctag acgctgaaga agagcggttn tgaattggcg 360
aacaccacgc ccacttgatt gcggatctag tctgtgagct agttcaa 407

<210> 49
<211> 420
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 49

agctntaaga cactagtctt cggccaaactn tgtttaaca ccaaggcaat taccatggtt 60
gctcataaat ccatatttac agaacaaaat tgnggcatgg ngctcaatac aaatcanaaa 120
gaggttctaa atatgttata gactaacaac ggcattcaat tagacaaaga gagacttagt 180
tctctaagaa tcaaattcgc atgcaaatttgg aaaattatag gatttggaaa atcatcacct 240
tttccccacct atcttactc ttcaaaaaccg aanatgattc caactcttct ctttcctta 300

gagagaaaata catgaagaaaa ggatggatga agattattcc tgcacccaaa cggagattct 360

aggagcttan naattcactc tttatnatat canaatacaa ggaatcttan aaattactac 420

<210> 50

<211> 423

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 50

tcatgttaa tcatacattc aatcttatta tagcaganaa ctcaaataaa aattcttatta 60

aaaggttagca ctcgtggcat gagtccttat aaatatctac tacaaaagtt ttaaatactc 120

tgtccatgag gaaagcagtc cagtatcttc caatactcta tccattgatc catatttgac 180

ctttcccat accctccata tccaatgcc aatggttcatc tccttcggga gcatcaaaaat 240

aagtctgcgc agtttacgga ttgacttcat cagttntcaa cctagtggtt tttttttcag 300

tctaagttag gacattacta tgcaagatga ctttgtccca tgtcaatggt tgagttcttt 360

gaacaactat tgctgactgc accacgcaac actatacgata tcatttggag aggcaacatt 420

cat 423

<210> 51

<211> 401

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 51

agcttgcttc tacaatagtt tctcttacc tcatactgaag agttacgaag gttccataga 60

agaagaaaagt ggctccaatt gcagaagaac cataaggtcg tcggttacat tattcattag 120

tcttggcgt gattgattga tgtaaagatc ttgcaataat cgtcgaanaa caagtagatt 180

aggagccata tacggattaa ggtatttcat ctaatctta ataatgaggc atgttgtaaa 240

tccttagggct tttggtagat tggtcttagt tacgcacatg ttgaattnta gcttccgcat 300

aaagaataaa gaatacggat canaagttaa aattctaaca actataagat gaacatcagt 360

tgc当地 attc tttgcacatg cattggatnt aaagaacaaa t 401

<210> 52
 <211> 464
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 52

 tagactaaat tcagcctacc atcctcagac tggccaa actgaacgga ccatttagtc 60
 attggaggac ctttgaagg cgtgtgtctt agagaaaaa ggaagttggg agagtttct 120
 tccattaata gagttcactt ataacaacaa ttttcaactt acgattcata tggctcctta 180
 tgaagctttg tatggtagaa ggtgttaggac acccctatgt tggtaaagc ccggagaagg 240
 ccttacctta ggaccggaag tggtaacaaca aaccaccgag aaagtcaagt taatccagga 300
 aaggatgagg actgctcaga gtangaaaa aagtttatcat gataagagga ggaaagatct 360
 gaaatttgag gttggtgatc atgtattctt gagaatcaact ccgtggactg gggttggtcg 420
 agcattgaaa tcccgaaagc tcacacactca ctttatcgat cttt 464

 <210> 53
 <211> 342
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 53

 agcttttagga taactttata ttngggaat gacagtagac ccacaattgt atttcatcca 60
 cagaactcta acccacaatt gtatttcatt cacagaactc taacccacaa catactagga 120
 tcttcaatca tatatcatcc caacatgaat aaataagact gattaagagt cctcaagctc 180
 ttaaatccaa gcccccccttg atctttggat ttacaaatca tctaagaaat aagatgaggt 240
 gttctgttat ttgcattcaact tcctgagata aagtctctgc agaagcttc aatctcatta 300
 cgaatagcaa tcgtaataaa ggttgcttca agaacataag ta 342

 <210> 54
 <211> 460
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 54

ntnttattgat tnnntttatg tcttgaccaa gaatttagaga ctaattcttt gaaatattga 60
gatttattta aaagataata tctcccaaaa aaatgttatt atattcaaac tgattaatat 120
gaacatgaaa aataaacaaa acgccttata aaattttagat ttagggaaatt attttaatt 180
actttaggaa atgttattga tgcacattc aaaaagtatg tgaaaggatg agtgataaat 240
cataaaattt ggctgctata agttatatcg ataagattaa atttaatttt taactcaaga 300
attaaggaaa gcttcataa aaagagaaaa atcaaatttt cattgacat gataatggtt 360
agagcctaaa aataaaatat aaattaaaaa tatacatatc aaaatacatc taaattaatt 420
aaataaaaag tactaaattc attggaaact agaaaatgga 460

<210> 55
<211> 371
<212> DNA
<213> Glycine max

<400> 55

agcttgtaca ctacaacacc aacaaagtcc aagaatccct ccataacaat gatgctcaat 60
accaacacgc tcttcccac cttctctcg tctctcaggt atatttgc当地 ttcatgc当地 120
ttgatatgct catatgcaaa aactagttc aaattttatt cttgc当地 tggtttgtt 180
attatatgca tagttgtca atcttc当地 aaactttatt ttaatattaa tggtatgtat 240
tgaatgttt taatgggtga gataggttagc actgacacag aagtgc当地 ttattggca 300
gttggaaagga gaagagatac ttgagcaatt cgaagcttct agttctctg agccggc当地 360
ttctataact c 371

<210> 56
<211> 459
<212> DNA
<213> Glycine max

<223> unsure at all n locations

<400> 56

ntgaaccatc cgccccatag gagcccaact ntgcaggatg aaggggtcaa acaaaaatttt 60
gatgaagcta caaaggagaa ctttggata ggaatggct taattgtaga aaccatcagc 120
gggagatcct aggggttgcc atgagcatgg tgaaccattg ttatc当地 cacatggcat 180
aagctgttagc tttcaaattgg gccttaaaga tagctaagga cctctttttt tttttttt 240

ttgacattgt catggaaaca gattgcttga agattgtta gacttggcac aacacaagga 300
agttttcaac ttcctatTTT gaaggcatcc ttGatgattg tagagagctg cagagttagag 360
gttttcatac gttcaaaatg tctttgtaa agcatacagg aaacaaagta tatggttcgt 420
tagtgaattt agctcttggt ttagggAAC gttattqqa 459

| | |
|-------|-------------|
| <210> | 57 |
| <211> | 423 |
| <212> | DNA |
| <213> | Glycine max |

<223> unsure at all n locations
<400> 57

| | |
|--|-----|
| agcttctaca ggtggtcagt aagaggaata gtgtaaggaa ttatggaaga taaaattcc | 60 |
| tagcaaaata acagttttg cttggggct aataaaggac agactaccaa caaggatgca | 120 |
| tctgcataagg agacaagtgc aactgcagga tctacgctgt ccttnttca gagaagctga | 180 |
| agaggagtca tctcatttg tcttccattg cgtcttcattc caaccaattt ggtgggaatc | 240 |
| gatgtcttgg ttgaatttac aaagtgcctt tcctcttggg cctaaacaaa atttctaca | 300 |
| gcataatttc atccaaggcag aaggttaag gattaagaga tggagatact ggtggatggc | 360 |
| agtaacttgg gccatttgga aattcagaaa cataattctg ttttcaaattt cagaatttga | 420 |
| tgc | 423 |

| | |
|-------|-------------|
| <210> | 58 |
| <211> | 467 |
| <212> | DNA |
| <213> | Glycine max |

<223> unsure at all n locations
<400> 58

tttgtgttag tagtgataac aagttcgta ttggtgacaa agagagggaa atcaaggaaa 60
gcactaacat ggaaaatgaa gtttgccaac gtgatgaata aatgatctag agcaaaaagg 120
cactgaaggt aaacgacatg gccccattcaa agaaggaaca taagtaggtg ctatcactag 180
tgcaaaaaaaaat gcatattaga tcaactttttt ggatcagctg tatgagcact gatTTTTTTT 240
aagaaatgca ataacaattt tgtaaatgaa aagaatgaaa tcggtctaaa aaatactacc 300
ttttggactc atggggta ttgtaaatgaa tttttgcattt gtattagaaaa catgagtcat 360

ataaaaacaa atgttgaatc gttntanaac aaatttacta tggaattntg tagcacatga 420
ttaacctang tcacactcag aaatatgatt caatatgacc cctacat 467

<210> 59
<211> 361
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 59

agtttccttg ataagctaga gcttagctac acacacccat ctaanaacta agtcaccc 60
cttgacaaaa tacatgagaa tacaaaaaaaaa aagtccctac tacaagact actcaaaatg 120
ccctgaaata caaggctaaa accccatact aatagaatgg cccaaataca aggcccaaaa 180
gaagggaaaaa cctattctaa tatttacaaa gaagagtgg a tccaaccc 240
tcaaaaaatct accctaagg tcatgagaac cctagggcct tcttttagtag ctctagccca 300
agcctttga agtcttctat ccaataccct tgggggttag gattgcata ttctgcata 360
g 361

<210> 60
<211> 450
<212> DNA
<213> Glycine max
<400> 60

tggatttcct ttttagtaggg aatctatcct tcctaagata gagccaaacc tagtcaccc 60
catataagaac tagctctttt cttcctctat tgccttagt tgaatacacc tttgtttgat 120
tctctatgg tttcttaacc ctctcatgca tcttctttac aaattctgac cttagattccc 180
cttctttatg tataaaaagaa gtgtccagtgg gggggaaat gaggtctaac ggtgttaggg 240
gattgaaccc atagacaacc tcaaaagggg actgcttggt gggtctatga acccccctgt 300
tgttaggcaa ttctacatga ggaagatact catccaaaga cttatggttgc ctttcagaa 360
gagcccttaa gaggggtggat aaagacctat tcactaccc tggggccca tcagtttgg 420
gatgacatgt ggttagagaac agaagtttag 450

<210> 61

<211> 247
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 61

agctttgtga atctcctgat ctagcgccac ccgatgatac tcaaccattc gaagntgaat 60
gtgatgctag tggaattggc attggagctg tcttgataca caacataatg cctatacgctt 120
atttctcgga gaaagtggga agagccttgc tgaattattt cacctatgac atagagatct 180
atgccattgc gagagcttctt gatcatttggaa atcatttattt tgcggctaattt cacttttat 240
tggattc 247

<210> 62
<211> 442
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 62

taccatcctc agactgaggg ccgagatgac atgaccattt agccactgca cgaccccttag 60
aacgcgcgtg tcttatacgcc tataggaggc tgagagagga ttcttcattt aatagagttt 120
acttataaca acaattctca ctctacgatt catatggctc cttatgaagc tttgtatgg 180
agaagggtta ggacacccct atgttggata aagccggag aaggcccttac cttagcaccg 240
gaagtggtaac aacaaaccac cgagaaagtc aagttaatcc aggacaggat gatgactgct 300
catagtaggc aaaaaaagtta tcattgataag aggagggaaat atctgaaattt tgagggttgtt 360
gatcatgtat tcttgagaat cactccgtgg actggggttt gtcgagcatn gaaatcccga 420
aagctacacc ttacttttatac ga 442

<210> 63
<211> 371
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 63

agcttatcga gaaaagaaaat tgtagaatgt ttgttaaaca cattgttaag ttcaactaaa 60
accctttgtta gaggcattttt cccaaatgtctt gtaagaccaa ctgtaaaaga aaaaaaaatta 120

aacacttgac aatggatgca tgcactacta tcactatacc agctagcttc attcgctctc 180
ttcaaggatc tatagcaatt ctttgcataa naatcttcaa actaacactt ggacagctag 240
atctaaccgt tgggtcgaa gtgtgacca attaatggtt atatattatta tgaataattg 300
aatattanaa tactcttggc agtgcatacc tacanagctc acttggggaa caaaacatta 360
cggtcttaa t 371

<210> 64
<211> 442
<212> DNA
<213> Glycine max

<400> 64

gtgcttccac aaaatagtct cggccgaaag acgctgacat ctccggaaa ggtgcagatg 60
accacattgg tctctgcgtg tcatcgact tgggtctcc gaataacgag gtgcggataa 120
ccgtaaagtg ctctgcatgc catcgaactc ttgggtcgct ggatagcaag aaggtgacac 180
taaatagtct cagtcgaaag acgctcacag ctccaggaag agtgcagatt accacattgg 240
tctctacgtg tcattggact tgggtgtcc gaatgatgag gtgctaataa ccgtaaagg 300
tctccgcatt ccacccgact cttggccgc tggatagcaa aattgtgaga caaaaattgt 360
ctcgaccgga agatgctgac atctctgtca agggtgcaga tgaccacatt ggtctccatg 420
tttcatcaga cttggatct cc 442

<210> 65
<211> 376
<212> DNA
<213> Glycine max

<400> 65

aaaattacta catatcatcg tatgcttata cctacacgca ttttgcac ac ctccaattgg 60
ctctatacg gaggtacaaa aactgcatca cgatcagctc actatacgaa aactataatt 120
caactcaatc ctcgccaatg tgtatggatg atctacgag gagcatgaaa cagacctcaa 180
cacacttgca tgatacttag agaacagtgt gttgcgaaa aagtctgcgc tatacttgcg 240
ccttaaccac aaggcgctat atccactgaa cctgtactgg ttagaccacc acctacggct 300
cgtagctcac acaacttag agcacctatg ttgagtccta tccaccata acgcacgctt 360

aagaatatct taagcc

376

<210> 66
<211> 370
<212> DNA
<213> Glycine max

<400> 66

tgaggatga ttgggttta ctttatgctg atgacgctac acaggatagc tttggataaa 60
gactgggctt acttgtacaa accatcgccg cgagatccta tgggttgcctt tgagcatgg 120
gaaccattga tatcgaacac acatggcata agctgttagt gtcacatggg ccttatacat 180
agctagcgac ctctttttt tttttttt tgacacttgc ttggaaacgt atcgcttgc 240
cattgttag acttggcaca acactaggaa gcttgaccc tcctattgtg aaagcctcc 300
tgacgattgt tgacagctgc acagtataag aattcatacg ctcaataagt gtttgctca 360
gcatacagga 370

<210> 67
<211> 378
<212> DNA
<213> Glycine max

<400> 67

agcttggca tagcaaatga gaaaaatgag tgacaaatgt gaaagcaaga gtcatttcta 60
ggtaaattt ggtgttgggg ggtcaaatct tgaatcggtt gagtttcgc cttacaatca 120
cttgagcaa gtctaaatta atgttatata ctcgtttgag atgagaattt actccaaaat 180
taccccatc tcattttcac ttctcaaacc ttgaaaattt actcaattaa tgggtttgg 240
ataccttagat ttggatttac cttgatctga agctggttt tgcgttaat acaattata 300
catgatttac gacttggtagg atccaatttgc agaaaaatg gatgtggca agaatggatt 360
cgaaatctgc cctattat 378.

<210> 68
<211> 296
<212> DNA
<213> Glycine max

<400> 68

ttatgttggg gcccacatgg atggtgcata aatgtataat cattatcgct atatgcata 60
cctggaaatg atttggggca ttcccttatt cctgaaccac ctgtgaaaca gacagccga 120
catacatcat gtctcgccac ttggaggcct tttgagccaa acattaactt ttggccataa 180
ccttggccta agatggaaat ttccaacctt accctccgaa gagagaacaa acgaatctc 240
ccaaacgaag cttctttac cttgagttat aagtgtcgag ccagacaacc gattag 296

<210> 69
<211> 365
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 69

atctntatgt tgngcaacac tgaaatacag actaaaaaag caaataaaaa aatatgctaa 60
aggcgactga agcaaaaaga gacaaaagtc ctccaaattt tacaaggaag gcatagaagt 120
gtaatgagga ttaatgtata agacaaatgg agtagagccc agcccaaata gttgaaatga 180
ataaaagtaca actaaggctc tcaaggntct tactcaatat aacccttaaa cactctntga 240
gcctttctga tcctttctt catagccttc gtacccctga ccacgttaca agcccaacaa 300
agcccatgtg gatcaaggaa ggactaatta tgctttgag tttggattct ggaatagaac 360
ccaca 365

<210> 70
<211> 321
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 70

nttcatatgc accaagaata gctcaataca cttatagaac tctcagatac aatgataaga 60
atgaactcat acttttgatg aaattccata ttgagtgtat tgtggaaacg acgcacttca 120
tgtgatgtgc attgagcaga tcttcattct acaacagatc ttcttatact ttagctattg 180
attacttctc atggcttaaa gttttactct tcatcagtga gcaatttgac ttcttcattg 240
cataacataa ccagagacac ttgcagccc tcttttaagc attctcgcca atgatcctt 300
ctgaatgact catgacaaca c 321

<210> 71
<211> 311
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 71

agctttaga attcacccca atttcgatga cctatgctaa ctgcgtccca aatctactcg 60
ataattcgat ggtagccata accccagcca aggttcctca acctccattt ttccgaggat 120
actacttgaa cacaacatgt gcttatcgtg gatgagttct agggcattcc attgaggcatt 180
gtangaccct gaagcataag gtgcaaggc taattgatgc gaggagaatc gcttgtgatt 240
tctgaaattt caagcgacac catacatggt gcaatttcaa gggtgttggt agatgtctct 300
aatgactcat t 311

<210> 72
<211> 423
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 72

nttcacanaa tcattaccaa agagttntac tctctgataa tctattacca gaaggtacta 60
atcgattact agtgttttaa aacattaaga tttcaaattt caagagttac aacttgtt 120
taaacatttt taacttgtgt aatcgattac acaatacttg taatcaagta ttgtgttatac 180
gattaccagt gtttctaaat gtttaattt tcaaaattca aatgaagag ttacatctgt 240
tgatgtgtgg taatcgatta ccagtgactg atttcgaaaa atacattcc aaaagtcaca 300
attactcaag tgacttgttt ctgaagattc tttcaaaaatc cacaactttt taagtgacta 360
gttnntaaaga aattgccaag agtcataaac tntgacttga gttatcaaga gattataagt 420
atg 423

<210> 73
<211> 397
<212> DNA
<213> Glycine max

<223> unsure at all n locations

<400> 73

agcttgccac ccagcttgc caggcgagca aggttgcttc ctccagaagc aacaaccttc 60
tggaggaatc ttctggaggg cccaagtggg ctttggtgct atttgcaccc ccattttac 120
taaatacacacc cctgcccttt tttggtgatt ctttttcgt aaagttacgg aaacttacga 180
atttcgtaac gatacttatt ttctttccgt aatgttacag aaccttgcgg attacataat 240
catcccccttt ttgacttacg gaatgttacg gaacctcact atttgcaca cgatgcttcc 300
ttttgatttc cggtgtgtca cggAACCTTA CGGATTGCAG ATCAATATT TCTTTGATT 360
tccgcacgac acgaaatttc acaaatngcc taatgat 397

<210> 74

<211> 465

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 74

tgttagaact atcatcacat .gacgctntat tggcacagaa catgttgctt tctaagcaac 60
ttgagattnt aacagaaaaca ctcggtaagc tgccaaactaa attgtctatt ggtcaaccta 120
cacactcttc tgtttgcag gttataagggtt gtaccatctg tggtagggct catgaaatgg 180
gccaatgtat tcccactaaa gaaaacactc aagaaattca ttatatggga aatcaacaac 240
gacaaaggta tactcaagga ggattttcag gcttccagca gggccctat aatcaacaag 300
gacagtggag gacacaccct gncaaccagt tcaacaaaga cttagatggg cttcaaaaca 360
gtccaatcca acaagggcct aacatattca agaggactac taagctggag gagaccttga 420
ctcagttntt gcaggttaaca atgtcaaatac atanaagcac tgagt 465

<210> 75

<211> 418

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 75

agcttataact caataagagg atcccgctgc gcatatcccc gtaacccac ggcttgctga 60
acaatattta gtgcctgcag atgttcttc cacaggcgat cgatattgct taagattagg 120

aaccgctctg cttcttcat caaggctgct gcttggcgccatccacaatatac ctatggggca 180
gcattgggtg taataaaaag gtaaattaa aaagaaataa atggtaaaga aaattatgtg 240
aaaagacaag gccactgaac caattcaaga cgactcaatt ttttagacatg acacatgatg 300
gtaaaaagtcc aaccttatat agcacaaagt tattacaagt tgcactgagg tttgataaat 360
ccctctcccc cttccgan aatgtataat caattacttg atttaaatca cttatctc 418

<210> 76
<211> 399
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 76

tgcctanaga ggtccaggaa ggacaaggca gccgaaggaa ctagttccgc tccggagtat 60
gacagtcacc gctttaggag cggtgtacac cagcagcgct tcgaggccat caagggatgg 120
tcgtttctcc gggagcgacg cgtccagctc agggacgacg aatatactga cttccaggag 180
gaaatagggc gccggcggtg ggcacactg gtactccca tggccaagtt tgacccagaa 240
atagtccttgc agtttatgc caatgcttgg ccaacagagg agggcgtgcg tgacatgaga 300
tcctgngtaa ggggtcagtg gatcccgtt gatgccgacg ctatggcca actcttagga 360
tatccgttgg tggtaaga gggccaggaa tgtgagtt 399

<210> 77
<211> 397
<212> DNA
<213> Glycine max

<400> 77

agcttgctgc tattcttgc tatgtgtact gagatatttt ctttgagctc tgatgcaaa 60
aatgatttat ttgcattttaa aacatagat ttaaccttaa atttcacccca aatcatagtt 120
ttcttagcaaa agttacaaat aaaataagtt taaggacctt tagtaaaatg aaaatatgcc 180
ccatatttgg actgagatg acaacagttt ggactattt tattaacgtt ttgacacctaa 240
aaatgagttt tctatgttttggaaaatgtatg gtagcgtata atatttgtga gaatccgact 300
aacagagcac caagagcact aaacataagg tatgagcgaa actgtgaaga actgagtcac 360
aaagagattc tattaccgtt gatgacttaa ctttgaa 397

<210> 78
 <211> 461
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 78

tgataaaagag attntgatag aagactntga ggccaaacta ttccaatata aaatgagatt 60
 ctttaacaa ttaactaag tccttatttg tgattttctt gatcttattt gttgacttgg 120
 aacaatgtgt gcttcatca agtaaattct tttggcatca tcaaaacctg cacgattcac 180
 atttatgtca ctcaacctct aggttttag agtaaaagga aggaattaat ggtgtacaag 240
 ttgaataagg ccttgtatgg tcttaacaa gcacctagag cctggaatag aaggattgac 300
 tcattttca tttagaatga attcaccaag tacacaatgg aatatgatgt atatgtgaaa 360
 aggaaaaacta aggaaatact tttgatttgc ctatatgctg atgatttgct tgtgactggc 420
 agcaatagtg aagaaataga gaaattcaaa gttgagatga t 461

<210> 79
 <211> 347
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 79

agtttcctt attctgtgac tcttcgttg gaaaatcctg ctgcatttc agattcgctg 60
 tgagttatca ctttatttgc ttctcctaattt ctctccttctt gttcacttcc tttgtttctt 120
 tggatgtttt caggatgact cctatgtcga cagctacgta agtactattt gagttgattt 180
 cgttaattttc acttttcctt tctttggta tttctattta tggcccaaca ctcattctt 240
 tttatatttcg naaatcaaaccatggcttg gaagggaaac cgccagctgc agattgttagt 300
 attctaagta gtggcggttgc cacgatcata tagcagtgtt ggacttt 347

<210> 80
 <211> 445
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations

<400> 80

tcagggtata aagaaaacttc atgaggcttt gttttaagc tataattcgt attgctcgct 60
tagcgcacag cccccccttat cgagtcata taacgattgg ttttaacaaa gccttgtgct 120
tagcccaacc tcgcgctaag cccaattcca aattttcaaa tcccagagag ttttggggct 180
tagtgagta ggcctgcgct tagcaactgtc tgcaactcaa aattttctg caatttgcgc 240
ttagcatgag atgtcaggct tagcgctaaa tcaagctcta acttacaggg atagtccang 300
cttagcgcatt ggaatgcgct aagcataatt ctatgagttt caaaaatagt gaaggattgg 360
cgcttagcgc atcttgcgc taagccaat tcatgaaaagt tcaattccag ggaggaaatt 420
gagcttagcg cangacagcg cgctt 445

<210> 81

<211> 387

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 81

tttcttatg attcattcta tgacccata gaggtccaca ttgtgtgttag agcattttta 60
ttctcggtgt gtgtactttt tataccccc gtcgacgtgc ttaagccatt ttacttaagt 120
cggttctcgc ttaacttaaa aataaaatag acttccaccg aacgtgtgaa ttgattcata 180
tctgacatca tattgagttt ataagcatct tcttggaaag ttgaacaagc ataagacatc 240
atattgtaaa tcaaacaagt actaaaacta tgcaaccatc cgtgttcat actttcagta 300
tcgtgtttaa attatgatgc atatcanatc atcatgaaat tcctccactt ttgaaagcac 360
caatgaaaatg ttctgtctcat ggtcagt 387

<210> 82

<211> 294

<212> DNA

<213> Glycine max

<400> 82

cccgccctcat gcctacgata gcaacactct aatacttcca taacgttagt agttatccat 60
aaacacctgct tagttgcct atgagcataa actagagtgt gatgctacat agatttaccc 120
tgccaccacgc cattagttag atgagtatgc gtaaccacat agaaccactg attcggaaaca 180

tgatatacat atgacgatga gttattctga tcctagatat aattaggatc cccgttgcac 240
atcggtggct catgcacttt cgacttagac accaactatg gatgagtcga ttac 294

<210> 83
<211> 315
<212> DNA
<213> Glycine max

<400> 83

agcttctatt attagctgaa ccattgtatc tatacacaca agctgagtgt tattcagacc 60
attagagttt atctctttta tcttagtgag agtgattctc ctaaattctt gagtgattca 120
agaacacccct ggctgtatca aaggacttgc acaacctttg tgtgttgcggc tcgctggaaa 180
gagtgattct ttccttccta tcatctccac ccttggctt tcaaaccaca attccagaaaa 240
atccacctct gccaaaatt atcttgac cataactccc atgttacaca ctcagattaa 300
gtgattcttg agcct 315

<210> 84
<211> 449
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 84

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caagatgcac ttggaaact atcgactgac ctggctattt tgcaacctgc acactctaatt 120
gattttagtgc tatacggcgc accatatgag gagaggctca tgacttgagc ctgtggatttt 180
tcactaacga taacactcaa gagattcatt atatggagg tctgctgcga ctccctgtatt 240
ctgccggatg attgttaggc gtccagcagg gtacctataa tcaacacaga cagtggagga 300
cacaccctct caaccagatc gacgaagact ataatggcc ttcactctgt ccactattac 360
aaggggctaa catactcaag aggactacta ggctggatga gaccttgact tcagtnntgc 420
agtagcaat ggcaaatacat aacagcact 449

<210> 85
<211> 361
<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 85

agtttcagac tgctcaattt ctccaggctg ctgcataggaa gggcaaaagggt ctgtatggtg 60
gtcagcagag gagcacaaac cacaaccct tgcgacaggt acagatttct gattcaaggc 120
caactgggtt accaagttga ccaacgcatac cagttgcct tcaagttct tagttcaga 180
tgatgcagat gggttttagt ctacccatg cactcctcta atgattatgg catcatttct 240
ggcgctaaac tgctngagt tggaggccat cttctcaatt aaatttctgg cttagcaag 300
agtcatgtct ccaaaggctc caccactggc agcatctatc atacttctct ccatattact 360
g 361

<210> 86

<211> 344

<212> DNA

<213> Glycine max

<400> 86

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gctctccaaa ctttggaaaac ttgcgttatt catctttca ttctcttctg cctttgccaa 120
aaagaattcg ccaaggacta accgcctgaa ctcttgatgt gcctctcttc tcctttatac 180
aaaagaacaa aggactaacc gcctgaattc ttttgttct cccttatgcc ttgacaaaga 240
actctgaacg acacaccctg agaattcttt tgattattgc cattccctaa tacaaaactg 300
tcaaaggact agccgcttga caattgtttt cgatccccat tcac 344

<210> 87

<211> 402

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 87

agcttaccat tataaagaag cttgtgagga agagttcca gccaaaccaa caagttctgt 60
tattcaactc aaggctcana ttgtttcatg gcaagatcaa gtctataaca gccgttaactt 120
ttaataaata aataaataaa taaataagta aaaataaaat aaaataatat aatttaggtca 180

taaatttcca ctatataaat caaatgttaa cctagagcag ctttacaaa acacttatgt 240
cccttctct tcttctgacg cacaagaatc ctaacagacg aactggagga ggagctctag 300
agagcaccag agacgccaca attgctaatt gagaacgatc gagggactac atcgaggtaa 360
gggatgagtt attcacgctt gnggattaga attaacatgt at 402

<210> 88
<211> 451
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 88

tgtatatcgat tacacacata ctgtatcgat ttaccatagg agttnttcag aanacattct 60
caatagtacat atcttttgt gtggttcttg aatggctatc ataggcctat atatatgtga 120
cttgagacac gaatttggaa agagttttc agaacaaaaa ggtctgatcc tcttataaag 180
caaaatcgcg ttatccttta acaaattcct tggccaaatt acttgtgatt caataaggaa 240
ttatggagt gctcaaattt ttcaatctat ctcttcaag agagattact tctttcttc 300
ttctttattt tgaagaggga ttaagagacc gagggctct tggtgtgaaa gaattctaaa 360
cacaaaggaa gggtagtcct tgtgttttta gaactcgatc aagaaattta caagatagt 420
gaactctcaa gcgggttgct tggggactgg a 451

<210> 89
<211> 563
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 89

gctcgccctca ctacacctcna cctntcatac tccgagactc ctatacatac gtagctgtaa 60
ctatagtacat acgactctgc tcctaaggct cgacccgcac nnnttgtttgcatgcata 120
catacaggcc attcgagtcg gcacccgtga ttctgtatctgcctgnat gcatgcaagc 180
ttgtttata ttctcaccac ttgagatgt gacaaccctt gtatctagaa acacataccc 240
atactcttc cctagtcgat cactcactta ataatacata ttctcgccct ttgatggtaa 300
gtttatgctt cacttcgaat tagatcaatt acttacgcga gtccttgatt taatccctat 360

ttccttcccc ctttggcatc gacataaagc acaaggcgac caccaatctt aaacatacat 420
acatgaccaa tcttgcaat atagtcgtt aaagaatatc ttgccgacta gcaaaggact 480
gcgtatgcac caaaatattt tccaacagtc attgagatga cggaattgtg gccatcatta 540
tagtgcaaaa gaactaatta tcg 563

<210> 90
<211> 559
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 90

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ctatcccccc tccgcncnc ttgattgatt ctgcgtggc gtgacctata catactcaag 120
ctctagcatg aagacagtac ctcagatgga gacttcatc ttgatgatac tacacaattc 180
gcgataggtt ttttataagc agagtggctc acgagaatct ggtcctaaat gtttcccaa 240
gcgcatgcaa tctctggaa tccattacca caatagtgtc gtgcgttact actagcagac 300
tgcgcttaac actgtttcgc actgaatgta cgtcgccct gtggacttac aatgagatgt 360
catgcataac gctgtgtcgg tctttcactt acagtgacag agaacgctt actcctgggt 420
cgctcgtaaa caggctata ctattccata gactctcggt gagacggact cactgcatcg 480
tggacgggta cccttgctgg ctatgaaatc ccatgagagg ttctccataga cgggtctgctg 540
ctttaaagtg ctctacatc 559

<210> 91
<211> 396
<212> DNA
<213> Glycine max

<400> 91

agcttgtaga attatgggt acccatcaca tgtggacta ggtggcggtc gggcgatgg 60
gcacaacaag ctttccacat acacaatgct cgcataaacc caccatcccc tggccac 120
ctccaaactga gtcacgtac tcccacgtt cccatatcct cgttctctc aacaccgggt 180
ccccatcaat cctctcaagc ttccacaaca tccaagcaaa acaacattca aacagcacaa 240
gctatcacag ccaagcaaaa cagagccaag gcagaaaaact ctgctcaaca catcaaccaa 300

aatcacagct tttctcacgt agagaccaca gtaacaattc ctgcgatcca attcgtaac 360
cgctggatcg actccaaaat tatactggaa gtctat 396

<210> 92
<211> 360
<212> DNA
<213> Glycine max

<400> 92

cttacgacca cgctctctcc actatttac tctcaaattga tctacattca ccatcccatt 60
tgtacttaca cccttcatt gtctatacac aagacacatt gatcttccac tggtgatgaa 120
tatgcaaggc tagacactcc atctatccaa ggagctactc caccactggc taaatatata 180
tatggtctat tcattgtact atctgcgaga gtggatcata ttcttgaatg ctatgtttga 240
taatcatgaa tatgaatatg ctgaccaatg ctaatgactc acgttatgga ttcatttgct 300
tcacttcgaa gatagacaca aagtgtttgg atgaactctc acctaatttg agatctcaat 360

<210> 93
<211> 407
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 93

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gagtaatttt gaaggaatga attggacta gaagaagaaa ttctttaagg atgctaattca 120
ttatgtgcgg gataatcctc acttgctcaa gattagagca aataaattga tataagatgt 180
gtcattttgg agcattctt ggcattatca caattcacct tatggnggcc attttatgg 240
agaaaagaagt gctgccaagg ttctccaagc angaattttg tggcccatgc tatataaaga 300
tgcacatttg tatgtgacac aatatgacaa atgccacaga aaatgaggaa tttcaagaac 360
gaatgagatg gccttgaaca acattcttga agntgaagtt ttgactg 407

<210> 94
<211> 287
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 94

tatgcgtgaa tctgggacct accatggcgg aagtctccac agaggcatt gcctccctcg 60
cccagaatta tgaccagtcg ttcaggtgct tcactttgt ggacttacag ctatcaccca 120
tggtggaaaga atctgaagag atcctatgtat gccctctagg gggaggaaa ccataacctct 180
tctcangatt ctatgcctct ttagctagaa tttctaagat agtccatatac tcggcgcagg 240
aattacacca cagaaagcaa gtcgaaaatg gagagagtgg agtaccg 287

<210> 95
<211> 380
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 95

agttcgaa taaaatgtatg aggtacaagc cctaaaggca gagcttgaaa gagccccgggt 60
agtcaaagag aagttcaagt ccatagccat canagtctga agagagtatg atgaactaag 120
ggacgtcaat atggccaccg ctgaaggcctt ggaatgagaa accatgaagg cccgaaagga 180
agaacatgac caaaacaaag ttntgagggg ctttataggg cagcaatagt gagctcaagc 240
tccaaagagg tgaaaggaat catcacggnt caaaggcatg atcttgaagg acgagctaaa 300
ggcttgcctt angtcgaaaa gaaatttgc ccaacagtta aagtgagact gaanggaata 360
tgtggccat catcgatgag 380

<210> 96
<211> 432
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 96

tgtaatcgat tgcacacata ctgtaatcga ttaccagagg agttnttcag aaaatattct 60
caacagtcac atctttgt gtggttcttg aatggctatc ataggcctat atatatgtga 120
cttgagacat gaatttgata agagtttc agaacaaaaa ggtcttatcc tcttataaag 180
caaaatcggtt ttatcctt acaaattcct tggccaaatt acttgtgatt caataaggaa 240
ttatattgagt gctcaaattg ttcaatctat ctcttcaag agagatttct tctttcttc 300

ttcttcattc taaaaaggga ttaagagacc gagggtctct tgttgtaaa gaattctaaa 360
cacacaggaa gggttgcct tgtgtgtta gaacttgtaa aaggaattta caagatagtg 420
gaactctcaa gc 432

<210> 97
<211> 370
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 97

agttgcaga ctataccccc gaccaaacac ggccgtgtt ctgtctcgcc ccggatttaa 60
agcgggttgc aacaccggct ccgcctccct aactgtactg gaggcggnng ccgtggcttt 120
gtcctctatg gtntctgga gtttaacat gacctccgag atgaaagcca tttgatctt 180
taaggccat agatcgacct tcattctgttc ctgctcgccc tcttcattat ccantttct 240
ggatttagtg ttatagggtt gccttgggt tttcttagtt atgatgaaat tcctaaagaa 300
ataaacaacg gtgagtatgc caccaaaaca ttagtatgca aatggatgat cggagcactt 360
ggatccaccc 370

<210> 98
<211> 458
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 98

tcatcatgaa tcaacaatga ttcanaggtt ttntgatgat aacaatgatg acaacaaaag 60
atgatgacaa aggtgatgaa caaaaagctc aagtgaatca aagaacatct caagtaaatc 120
aagaacaagt caagagttca agaatcaagg agaattcacg actcaagaag aaagcctaga 180
atcaagaatc aagactctct caagaatcaa gatcaagatt caagactcaa gattcaagaa 240
tgaagaaaag actcaatcaa gataagtatt aaaaagtttt ttcaaaaactt tgaataacac 300
atgagttttt gacaaaacct ttaccanaga gttttactc tctggtaatc aattaccagt 360
agcaaaaatga gttgaaaaa agtttcaaa ctgaattac aacgttccaa atatttcaa 420
aaggctgtaa tcgattacaa tgtttggta atcgatta 458

<210> 99
 <211> 349
 <212> DNA
 <213> Glycine max

 <400> 99

agcttctata aaggttctta tggactgctt gtctcatcat tccagtaata ccttcaccca 60
 caatattgaa aagcaaggga gctagggat ccccttgccc caaaccctta gaaggattaa 120
 actccttaga agggctacca tttatcaaaa ttgatatcga tgctgagtgg aggcaagctg 180
 atatccaaga tctccatTTT ggacaaaaac ccattctgca cagcatgtaa tccagaaaaag 240
 accaagaaac tgaatcgtag gcctttcaa agtccacctt aagaatcatc acaggtttct 300
 tatttctcct tgcttcctca accacttcat taaggatcag aataccatg 349

<210> 100
 <211> 432
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 100

tatcaaccaa atacccacgt tntcaaactc aaagtatatt acaattacaa aacataaata 60
 agaaaagatct tcacatctat aaaaagactt aatgaacatt aacagaaaata aaacctacca 120
 gaacctgtaa gtgaaagaaa atatacagac ccatgtaaac tatagaacat agatctaacc 180
 ttttaagtgg aaaataacat aaacgtagaa gtatgcaa at gaagatttac caaatgaatt 240
 cagtaatagc aactttacct gaaagtgtaa actgttaatg cagcacccaa tccgccaaaa 300
 caatgggacc atgatcctct ggaactctac catgttaata acttccaaaa tttcttccag 360
 ttcacccaga aacatcacct ctttctgact atttgttgct ggccaatatt tcaacaagcc 420
 gcttattcaca at 432

<210> 101
 <211> 406
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 101

agcttgcata ggcctaagg atggacaagt gttgttcagt gttgagagtg gtagccttga 60
caagtgttgtt cccgggttgct tctgtcaagt tgtctgggtt ggacaccctt tttgatgcaa 120
gaggctccaa gaggattggg cttagagctgc tgaagaaggtt cctaagggttc tcataactt 180
taggatagat tttttagcccc atggccaag gttgggtcca attatcttt tacgtttag 240
attacgatgt cactatattt gttcttgta attagggttc cataatgttag gttagggtacc 300
ctagaaatat aggattttc agcccttgta ttttagggca cctagacttag ttnttgatt 360
aaggtagtt ttgtatattt acatgcacta agtgaatatt taatgt 406

<210> 102
<211> 451
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 102

tntaaggccc tcaaatttactt ggtggcaac accccactta aatgattatg aaagagggat 60
aaatatgtga gatttttggaa caaattaaaa gaaggaagtg gacctgtaaa gttattggaa 120
gacaagtcta gataaacaag ttggctgagt tctgagaatg aacttaggaag tttccattt 180
aattggcagt aagcttagatc aattttagat aactgcttca tattggaaat tgcaccttgg 240
agctttcctg agaaaatttgt atagctaaga ttcatgtgat gaagagaacc atgttgggg 300
aagtttggca aagaaccccc aagatcttgg ttgtctgaga tgtcaaggac cttcaacgtt 360
gatattttggaa atatatcttt tggaaaagaa ccattcaaggc cacaacttct taactcttagt 420
gtgactaaat tggagaaatt acaaaggatt c 451

<210> 103
<211> 410
<212> DNA
<213> Glycine max

<400> 103

agcttgcgtc tattccttgta tatgtgtact gagatatttt ctttgagctt tggatggaaa 60
aatgatttat ttgcatttta aaacatagat ttaacccctaa atttcacccca aatcatagtt 120
ttcttagcaaa agttacaaat aaaataagtt taaggacctt tagaaaaatg aaaatttgcc 180

ccaaatgg actgagagt gacaacagt ggactat tattaagggtt ttgacctaa 240
aatgagttt ttaggttg aaaatgtagg gtagcgtata atattgtga aaatccgact 300
aacagagcac caagagcact aaacataagt taggagcgaa actgttgaaa actgagtcac 360
aaagagattt tttaccgta gatgactcta acttggaaatc caagtctctg 410

<210> 104
<211> 460
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 104

tgataaagaa attntgatag aagactttga gcccaaacta ttccaataaa aatgagatt 60
cttttaacaa tttaactaag tccttatttgc tgattttctt gatcttattt gttgacttgg 120
aacaatgtgt gcttcatca agtaaattct tttggcatca tcaaaacctg ca^tgattcac 180
atttatgtca ctcaacctct aggtttgag atcaaaagga aggaattaat ggtgtacaag 240
ttgaataagg ctttgtatgg tcttaacaa gcacctagag cctggaatag aaggattgac 300
tcattttca tttagaatga attcaccaag tacacaatgg aatatgatgt atatgtgaaa 360
aggaaaaacta aggaataact tttgatttgc ctatatgctg atgatttgct tgtgactggc 420
agcaatagt aagaaataga gaaattcaaa gttgagatga 460

<210> 105
<211> 411
<212> DNA
<213> Glycine max
<400> 105

agtttcttca gaccatgata ctctaggacc tcagggaaagc aaaatgttat gaattcaact 60
tggttgaatc aactcaatttgc ttagatagtt gtgccggat aatgcttcat agtgccttt 120
tgcaaaccat ggcataaatt tgggagggga tgagtgtttt ggagattccc cctttgtaga 180
tcacccaaaca actttcttcc tcttcttctt cattttctcc tctatgagct ttgtttctt 240
ctcttttaa ggcttaaggt taaagggagc attgttgatt gcaaccctct taatatgttt 300
ccttcttaggt tggtctaaca tggtgtatgg gaagaagaaa gtgatgggtt gaaggaatta 360
cggaagaaga aaggatcg aaaaagggtt acttagcatt cccaaaaact t 411

<210> 106
<211> 463
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 106

ttgatctacc accacccgccg ccaccatcat tntagttntc tattatcaa ttttactagt 60
actttgtttt ctagccgtgt atttggttat attatgacat ttggataatt tagtatttct 120
ttatggcat ggtttgattt aacaattatg aattatgtta tatgactatg tggttttat 180
atatttgate tattcatgtt tcttccttca tgattggctt atattcttca atgtatgtct 240
tgtgaatgt taatagtata tgtttgcct atactgtta cgcaacttgg cttttgg 300
atgccaagg gggagagaaa tagggattaa atcaagaact cacataagta attaacttaa 360
tttcaagtga agcatanact caaaaaacac aggccggagaa tttaagttag tgttcgacta 420
ggacaaaatg tgtgtatgaa tttcttgatt tcagggttat cat 463

<210> 107
<211> 406
<212> DNA
<213> Glycine max

<400> 107

agcttgcatt attgatggag aaaagggAAC aaccatgaat ggacaaagta atttagctgc 60
tactaactct gttttcact taagcaaATC aatttcttagt gatTTTAA catcgaaaat 120
gaatcttata aaattactgc ttcatcatca ttttagtggc ctatTTGATG gcaataacta 180
aaaacattaa cgaaatataa gtttaaaAGA ctaatcacAA ttttggg gaactaaaa 240
taaaaaacctg aagaaatCCA gacataataa acataatcta ccccaactta tccaaaatac 300
attacattct atttctata atccctttc ctTTTTAA gatTTTTA aaaatattat 360
attacaaaaa gtatataattg ataaaggcatt ctTTTTTC aatttt 406

<210> 108
<211> 388
<212> DNA
<213> Glycine max

<400> 108

tcgcgatgtat ggtcatgtatg actacggggg atgacgactc ctgtgaggaa cacacagctc 60
tactgagtca gagaacatct caagttattc tggatcatgt cttagagttcg tgattcaagg 120
agaattcgcg tgtccagaag aaagcctaga ctcagaatc tagagtctct caagaatcaa 180
gatcagcatt catgactcat gattcttagaa ttagtgaatag actctttccc gatcagtatt 240
aacgagtttt tgtcgaacct tgaatagcac atgagtgtt gacagaacct ttaccagcgt 300
agtttgactc tatggcgctc aattaccagc agcacaatga gctcgaaaaa agttttcaga 360
ctgaatttac aacgctccaa atatttc 388

<210> 109

<211> 318

<212> DNA

<213> Glycine max

<400> 109

agcttatatg caatgtggta ccatgtcagt gaataacctc gtcggcgcc taggagtaca 60
tgacaagaca aaccacacaa taagtagtca agtcaactctc actaggtaat atcataggga 120
gaccagtcag ggtcacagtg ttttgcgaga atgatccaac catatggat caacataggc 180
ttaaaggagc actcaaaccg tgtaaccccc aaggcctaca ctccgaagag ttcgtcaggg 240
cctctccctc ctgattcagg tccaaaccag aaaaattta gcacacagac tctatctatg 300
aactgtacaa aacacact 318

<210> 110

<211> 165

<212> DNA

<213> Glycine max

<400> 110

tcttatccaa cgctcatctt ggtggagaag ctccttcttc catggcttat tccctagtgg 60
atggtgtcctc ctctcacctg atctactttg cttccgcta tatctccatg gcgaaaaatc 120
gccattaaag gacctcatcg aagctcatgg aaccatccta catag 165

<210> 111

<211> 394

<212> DNA

<213> Glycine max

<223> unsure at all n locations
<400> 111

tataaagctg tttctgggtt tatttaagtc ctaagctata gcttccttcc tcattgtacc 60
ctgttcagac ttgttatata tatatatata tatatatata tatatatata tatatatata 120
tatatatata tatatatata tatatgggtt tgctaattgca ctcacaccat ttttttagta 180
tgagtacatt agcaatcaac accaaanatt ttaactaaaa aatcaatcac ctctcttatt 240
tttcaaaatt taatgttaggg atatcctaca tttgttccaa gctcgacaca tctgatctat 300
atgctcgagc ttgaggggga gtgtgaaat atgataagtt cccattggaa atgatctgtt 360
cctaattcta ctactgttcc aatgaaatta ctcg 394

<210> 112

*<211> 330

<212> DNA

<213> Glycine max

<400> 112

tagctggtcc ttgttgctt cttcacacac ctctctctct ctctctctca tgcactccat 60
atatcggtac ctagatacac aaacaaagaa aggctatac catatgaggg atatgaatga 120
aagaaagtct atagcgcagt gtcattttt tggattagaa atattcccc aagctgaacg 180
agaagccgc catgccacgg ctgtagatat ccgtttaat atgaaaacat tcccgtgggg 240
ctttcataag aacaatgagc cttcataat ctatgagaga aagcacaact catcgaagct 300
cgatcattac ctatactatc ttgtttat 330

<210> 113

<211> 229

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 113

gctttgatgc cagctatgag gtatgtcat gtttgaacag acatttatgc ttgcctata 60
ccccccctcct tggctcttag cgcaagttac tccttgaca agtactggga gctggagtaa 120
aatgatgaca tgcccccttag ctggactgac agcgactaca gtctggacta ttttagtca 180

cgacncgacc tcttaaagga cctttatatg attgtaaatg caggtagc

229

<210> 114

<211> 407

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 114

atgatataag actttgaggc caaactatat ccaatatcan aatgaggaat cttaaacaa 60

ttaactaag tccttatttg tgattttctt gatcttattt ggtgacttgg aacaatgtgt 120

gtttcatca cgtaaattct tttggcctca tcagaacctg cacgattcac atttatgtca 180

ctcctcctct aggttgtgag atcacaagga gggatttaaa ggcgtacacg tcgaataaaag 240

ccttgatgg tcttagacca gcacctacag cctggaatat aaggattgac tcagtttaca 300

ttgacaatga ctccaccaac tacacagtgg aatatgtgt gtatgagaaa aggataacta 360

cggaaatact tttgatctgc ctatatgctt atgatttgct tgtgact 407

<210> 115

<211> 378

<212> DNA

<213> Glycine max

<400> 115

agcttataacc agccaaatcc cccaaatttt ttaatccgag ctggaaattc tctcccgact 60

caagtaaaag gaccacctgc aacagaaaaga gcgcggccc aacgcacggc tccagccgct 120

ccccggccag ttaataatac agccccgac gcgacctata aatatgcaca gcacccgccc 180

ccgaaagata acttctcccc tattcccatg gcatactccg agttatggcc ttcattattg 240

gagaatcatt tgggggtggc cataccggg aaggtcttcc agccacccta ccccaagtgg 300

tacgaccgg gtgccaagtg tgtgtaccat agtggagctc ccggacaccaa tattgactcc 360

tgcatccgt tcaagtat 378

<210> 116

<211> 457

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 116

tccattgttg agttgttgct tccctgtca cgctctaatt cactccccac aagtaagtgc 60
aattccctt gtttatttgg ctctccattg atgtgttttg gtgcttagt tgctcatttt 120
tttgcaaaat tcgtgaagca attcgcatct gaatccatgc ttgtttgtt gaattgaggg 180
tttgtgtgag aaggcattat gcctatgttg tattctgaag caatggggca tgccacattg 240
tccccattct ctgcaattt gagtcgaac gtgcgcggcc caagtgctcg gtgaagtgcc 300
ccaatgatat atgaatatga tttgcacaa ttggatggt gggactgttt tatatatgtt 360
gagacagcat aagagattca aaatatgtgc ccgaatgcaa tttcaagctt atgaacccan 420
acctttatc ttcaatgcaa gaagacatac tcatacg 457

<210> 117

<211> 305

<212> DNA

<213> Glycine max

<400> 117

agctttgata gtaagtttaa ttgaataaat tatactcact atcacaaaaa tggtcttcta 60
cgacgcacgt ttacgacgg ttgtacaaaa accgatgtca taagtaaagt agtgacattt 120
ttgtaaataa cttaaaaatt ttaaagatgg ttcttatcaa accagtcttt gaaaaggaat 180
taccacatca gttcttctac aaccgacgta gaatgcgaag cttaaaaatg cgaacgggct 240
ctctctcact ctctattata tcctctttat aatctctctc ctccctaaaat cttagaaaacc 300
ctaatt 305

<210> 118

<211> 427

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 118

tctctgcctc ggntgaaaca ttggcgggtga catgtgacct catgtggcca cccaaggatc 60
ttccacaagg gaagctcttg ctgcagaact tgcacatgtg tttcacttct tgatcttctt 120
ccattggatt gatTTTCAA gatcacacac acgcacggat cagcaaagaa agcaaaatta 180
accacacttt cttgatcacc accaacacaa gagaatcga tcacaaggaa aaaacagcaa 240

caccccatcaca tcttggaaagt gggtggagag aagaataata ccgagaagaa 300
gaagaagaga aaccccatgt ctgaaaattg caaggtggtg agtgcaagat ctaacgcaga 360
aaacaagagg aaaagaaaaag ggacaagaga acgtgttagta gtagcacaaa ctattatata 420
tactata 427

<210> 119
<211> 394
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 119

agctntttgg agtagaaaca taggaccaac tcatttattt tcaaaaagaa agtcgtatct 60
agtcaaggc tgagagacca tacaagtttc ctaacgattt ctaattatgt gggccattaa 120
gtctatcata tgctgacaat agccgagaag cccatgaatc tttcgaaaaa cggagtaagt 180
gtctgccatc gccttggcct tggctaacaa gcggngaagt tcttgactcc cggtcaagg 240
aagagcaaac cgatccatcc acatggttgc ctcttggtgt aaagagtcga tcacccttcc 300
tctagcctct tttccgcgt atacttgagc atactcgtcc gcgattctat gctcgtggc 360
cgtggctaga cctaactctt cttggactt ggcg 394

<210> 120
<211> 335
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 120

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acaataacat catttcttgc tctgaattgt tggagttgg aagccatctt ctcaatcaa 120
ttccttagcct caacaggggt catatcacca agagctctac cactggcagc atcaatcata 180
ctcctctcca tggactaag tccctcatag aaatattgaa gaaagagttg ctcagaaatc 240
tggtggtgag gacaacttgc acacaatttc ttgaatctt cccagttactc atacaagctc 300
tctccactaa gttgcctgat gcctgaaatg tcttt 335

<210> 121
<211> 439
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 121

ntaacctcat cgtctctcac agtcttaga attgggagcc aatccaatcc ttgtgttcgg 60
actctcagcc acttatgata gccgtcgatg atcccattac tgctccctt aagctctatg 120
tcctttcttc atgccgcac ccatgcctt cgaactcctt ggagtaccct cacgttgtgg 180
tcaccgaaac cccgtgcgt gaaaggcgtg atgcttcgt ctgatggcac tcctctcatg 240
ggtagccaa gctgtcttat ggtgaggacg ggattataat taatacaacc ccttggttcca 300
tcaagggAAC atttggacat ctttcgcac aagatagaat cctgattctt ctttccttct 360
agcgaggAA caaattaata gacgcccctc catgctagcc aagagttggc cccaaattcgc 420
ctttcccttt tcgatgcac 439

<210> 122
<211> 397
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 122

agctttaAGC caattcatac'gacaataact ttntactcgg atgtctgatt gagtcccgta 60
atataacgaa acgctcgaaa ttgaatgtt aagctctaAC cctattcaAA caacaataAC 120
gttttactcg gatgtctgaa tgagtctcgat aatatATCGA cacgctcgAA attGAATGTT 180
gaagctctAA gcctattCAA ACAACAATAA cgTTTACTC ggatgtccGA ttgagtgacG 240
taatatacG agacggtcGA aattGAATGG tgaacctatG agccaattTA aacgacaATA 300
acttttact cggatgtctG attgagtccc gtatatacG agacgctAA aatGAATGTT 360
gacctctgag ccattcaaga caatactttt actcgat 397

<210> 123
<211> 427
<212> DNA
<213> Glycine max

<400> 123

taaacattca atttcgagcg tctccatata ttacgcgact caatcagaca tctgagtaaa 60
aagttattgt cgtttgaatt ggctcagagg ttcaacattc aattttgagc atctcgatat 120
attacgggac tcaatcagac atccgagtaa aaagttattg tcgtttgaaa tggctcagag 180
cttcaacatt caatttcgag cgtctcgata tattacggga ctcaatcaga catccgagta 240
aaaagatatt gtcgttgaa ttggctcaga ggttcaacat ataatttggc gcgtctcgat 300
atattatggg actaaatcag acatccgagt aaaaagttat tgtcgtgtga attggctcat 360
aggttgaaca ttcaatttcg agcgtctcgat tatattatgg gactcaatca gacatccgag 420
taaaaaag 427

| | |
|-------|-------------|
| <210> | 124 |
| <211> | 413 |
| <212> | DNA |
| <213> | Glycine max |

<223> unsure at all n locations
<400> 124

| | |
|--|-----|
| agcttggttc aaccctataa tccaaggaaat ggcaattcta atcgccaata cttcaacaac | 60 |
| atctcatagg gatgaatgac tcaggcatac tttaagctta tgcacggaaa atgtaattat | 120 |
| gaaattgaga tgcccgaga aacaccattt cctagttAAC catgcattAG gtaccatgtt | 180 |
| caattatttct gttttgttgg tgtgtgtttt tttttttttt agaaatgggt ttatgatccc | 240 |
| aacatggttg gctcatggtg cctaacacat gcaactaaga atgttagtgtg aagtttcacg | 300 |
| cttcccccttt ttgtttntg ttttgttagag gaaaacgcaa ggatgagcaa acatganaac | 360 |
| aaatggtatg caatTTTgca gatcanaaag tttgttgaac gcataATGcat gat | 413 |

| | |
|-------|-------------|
| <210> | 125 |
| <211> | 333 |
| <212> | DNA |
| <213> | Glycine max |
| . | |
| <400> | 125 |

tcaaaccttag gccatcattt ctgctccaaa tcgcgaaagg agagcattct tggagtgcgtg 60
aagtgcgtgg ctacgagtgg gacttcgaaa attcaggttt gggtggactt ctttctcctt 120
taatttcgt gggtatgggg tttggggaga tatgatgggt agtcttgcta ggtttctqct 180

gtgtgatgat tatttgaa gacattgtt gaaagcttgt tgaaattgcc atgtttggat 240
gagttagaca tacccattct gtttaggtt tttgtatgc atgctgtga tgtttatatg 300
ctgaaattgc ccatggaaaa ctgctagaga tga 333

<210> 126
<211> 405
<212> DNA
<213> Glycine max

<400> 126

agcttccttag ttccatgtt tccagatggg ttttagcta cctcatgcac tcctctaattg 60
actatggcat catttctggc gctaaactgc tggagttgg aggcattttt ctcaattaaa 120
tttctggctt cagcaggagt catgtctcca agggctccac cactggcagc atctatcata 180
cttctctcca tattactgag tccttcataa aaatattgga gaagaagctg ttctgaaatc 240
tgatggtggg ggcaactggc acatagtttc ttatatctct cccagtaatc atacaggctc 300
tctccactga gttgtctaat acctgagata ttcttcctga tggctgttgtt cctggaagca 360
ggaaatttt tttctaaagaa tactcttta aggtcatccc agctc 405

<210> 127
<211> 388
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 127

agcttgctct atatttacat tgatgcttat gggaaagaggt tggatgccat tttgtttta 60
agagtagtgtt cccactggta aaactaactt tccaaatgtt tgccttcgca gggaaatggcc 120
ccaaaggaaac ttgcctcata gaggtctagg aaggacaagg cagccgaagg aactagttcc 180
gctccggagt atgacagtca ccgttttaag agcgccgtac accagcagcg cttcgaggcc 240
atcaaggat ggtcgttctt ccgggagcga cgcgccagc tcaagacgtt aaagaagcgc 300
tactaggagg caaccttagta cttttaaat ttctgcctgc tatttgatca ctctttatag 360
tangacgcac ctaggtgctc atgatcct 388

<210> 128
<211> 458

<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 128

tgccctaatta acctagaatt gagaganaat gattattaaa cacaataat gaaaataata 60
agtatttatt acctataactt aacagaaaaat acttataaca ttacaaaata accataaatt 120
gggagagttt gatacaattt atacaagttt tatacacaaa agtttagtcat tttcaccgac 180
taacaactcc cccaaattta cagtttgct tgtcctcaag caaaaagaga acaactcact 240
agtgcgtcgag tgacaatgac atgcagtgac tatgtacaaa ggtgtatgct acaaagtgac 300
tgattgcattg ataagagaat ggagtaaaat gccctaattca cttgtcttc acaaggtatg 360
cagttatcca aagagaagaa taaattgtaa cctgaacaga tagatgaagt taggaataag 420
acagatataca aggaaagtag cttacaccat agtctcat 458

<210> 129
<211> 347
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 129

agctttacta ttatcttccg aacaatataat gaatgaaatg atgaacctta taattaaaca 60
aagataactac tactactaag tttctattga tgcttgatc tgagtactaa aaagaaagcc 120
tgttataatg attcaaaggc ataacaataa acaacttaac aataaaccat gaactacagc 180
agctggnggt actttaataa atctctttgt atttaaaat agtctctaaa attntatgta 240
aaaaagataa cttaacttat atttactaac taatgatata aaactaattt gctaacgatt 300
taaactaaaa caaaaatgcg ggaccttaat aaatctctt gattat 347

<210> 130
<211> 398
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 130

ntaacctcat cgtctctcac agtctttaga tttgggagcc aattcaatcc ttgtgttcgg 60

actctcagcc acttatgata ggcgtcgatg atcccattac tgcttccct aagctctatg 120
tcctttcttc atgccgcata ccatgccttg cgaactcctt ggagtaccct cacgttgtgg 180
tcaccgaaac cccgtgcgat gaaaggcgtg atgcttcgt ctgatggcac tcctctcatg 240
ggtagccaa gctgtcttat ggtgaggacg ggattataat taatacaacc cttgttcca 300
tcaaggaaac atttggacat cttcgcatg aagatagaat cctgattctt cttgtttct 360
agcgagggaa caaattaata gacgcccctc catgctag 398

<210> 131
<211> 312
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 131

agcttgtatg ggtgagttat aaattataat acctcagctt gttcaaaag atcatgttta 60
ctccgtcgca cgtttagcaa gtttgacaca gtttgcaca ctatggacat cttcacccga 120
ctagctctta ctctggcctg aactcgaacc aatgcctgca tgcaccta tgcacccgt 180
gtctgcttcc tcacctgtct ccccccgaac aagtgttga atcctcacca ctggcttcaa 240
tgccctcaaa gcccttcttg cctgcaatca ccacacaagt gatgtgaaa tataatgtt 300
gtctaattgtt aa 312

<210> 132
<211> 414
<212> DNA
<213> Glycine max

<400> 132

tgttcatgtg agagctcaac actaagttgc tcatgtctt acaaaggccct taatgcttct 60
aactctcttcc tcctttatcc acaagttggg actcattaac atttactctc caacttgagg 120
gggtattaaa gttgtatgaa gaaatggagt tagttacttc agatagaggg tagatagact 180
agttggtaat tagttagaat gaagtttagat actaagtttgc ttaagctgaa tataaaatag 240
tgtgtatgca accttatatt caataatcat caataatatt ctacagattt cttgttgca 300
caaagctctc tatcaataaa ttcccccttg ccaagtccac attgaagaat ttagagcaat 360
tgtagaatgt cgaagaacat attatgtaca tacaagacac aacttataaa tctc 414

<210> 133
<211> 397
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 133

agcttgctct aaatttacat tcatgtttgt atttatggga ggagggttgta tgccatTTT 60
gttttaagag tagtgtccca ctggtaaaac taactttcca aatgttgcc ttgcgaggaa 120
atggccccga ggaagcttgc ctcanaGAGG tccaggaagg acaagacagc cgaaggaact 180
agttcccgctc cgaggtatga cagtcaccgc tttaggagcg ctgtacacca gcagcgcttc 240
gaggccatca agggatggtc gtttctccgg gagcgacgCG tccagctcg ggacgacgag 300
tatactgatt tccaggagga aataaggcgc cgacgggtgg catcaCTGGT tactcccattg 360
gccaaGTTTg atccagaaat agtccttgag tttttagt 397

<210> 134
<211> 450
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 134

ctgatagcag atgatATGAT ctttactagg agtggatcgc ttgatacagg tcataGAGTT 60
tggatGATGC tacttccaga gagggaaGAT aagtcaGGAT agacaccaca agaattgcct 120
tgataAGTCT gagattgggt caacatgaga cccagagaga agctctctcc aaagtttata 180
aaaggccaaa agtacttata ttgaaaatga aaccataca tatagcgtat ctgaatgaaa 240
aaaatataaa tagaccaggg cttcanata agttagggcc aaaattacga caataaaatt 300
ataaataaca aatagaacat atttgcATG ggcttcaaa ttGTTGGG cttcaacaa 360
caattaatat tcttagtagt gcctctggct ttggaccttc atccttctcc acttgagccg 420
tggtaagtat gtctgttacc aatttGTTGA 450

<210> 135
<211> 394
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 135

agcttcttga tatactgcag cacgattata ttaatgcttc atagtatTTT ttgacatata 60
tattaatatt gtcaagctaa aggcacaatt tttgttatgc taattgataa caatgataga 120
aatgatctta gatacaccaa gctgccttgg atatgatgtt tgattcgatg atgattgggt 180
agttagttagt tagagaggaa gaattaacta agttaattat attaaagatt 240
aataatttga aatgagacta taaaacata tagataataa attgtcataa tgaaaagttt 300
acgatctgtt tggatatga taaaatagga taagatatac cattatgata aacttttaggt 360
ttatTTAATA ttgtatgcac actanataat atta 394

<210> 136
<211> 440
<212> DNA
<213> Glycine max

<400> 136

ctataaggac atgcatccag tcaacacacc gctgccaatt cccttatgaa tatcctgaat 60
cataaataag ttcttcaaAC aatagttAA gttatgactg aagtttcata tacgcaaagg 120
caaataataa taatagtaat tgagctaAGT ttttggAAA gtagttcaAA ttAAAataa 180
aaaatattaa atccatatta aattcgtaaa tttgaaACat tgtaattgta ctcaagtga 240
cgatTTTTT tttttttgc agcaggtcaa atcaaggctg gtatctttat tttgagccaa 300
aatttcttaa gcagcgTTTC acccgcaaAT tatgcaacgg cgaaccaAAA cttagaact 360
gaaaatcttg cattCCAAGG aaccCATTAG ccattggcCT caagacaAGA actacgtatt 420
gtggctgggt tgatcccTAC 440

<210> 137
<211> 340
<212> DNA
<213> Glycine max

<400> 137

agcttcAGC tatgtatCTC attagtaata ctTTTcGTG ctgttggatg ggcaacaatt 60
gcctcCTTGG tggtagtAgT aatcactgtg cttgcaata ctccactcgc aaagttacag 120

cacaagttc aaagcaaact tatggtgaca caagatgata gattgaaggc ttgttcttag 180
gctcttgta atatgaaggt gttgaagttg tatgcgtggg aaaccaattt tagaagttct 240
atagagagat taaggaatga ggagctcaaa tgggtgtctg cagtcaattt aagaaaggca 300
tacaacacctt ttctcttttgc gcctctccctg agttggctct 340

<210> 138
<211> 453
<212> DNA
<213> Glycine max

<400> 138

ctgcttgatg agaaagagggc acagttacgc caatttataa ggtaagcacg tgggtttta 60
ttcagtggtt tagcatctt tggacatgct tgggtttctg attaatttgg attgaagtat 120
ggaagggggg agtactggtt catcgagcaa tttgcgtgaa gaaactgctc gtgtgtcgaa 180
agattttca gagtttttc caactggcat tccacaagta ggtcaaacag agattagcca 240
agattcattt gccccgtggac tggggatat tcggtcgag ttgattggct ccacatctgg 300
caatgattct actacttttcaatcgaatga ccgtatgaga aatggcagag ctgacaatgc 360
caactctaaaa ggtcatgaca gtcatttttttggcagacag agatataactt cattgctgct 420
caccctcctt gtcaagcgat gttaattaat tgc 453

<210> 139
<211> 304
<212> DNA
<213> Glycine max

<400> 139

agcttgctct aaatttacat tgatgcttgcatttttggatggata tgccattttt 60
gttttaagag taatgtccca ctggtaaaac taactttccaaatgtttgcc ttccgcaggaa 120
tggcccccggag gaagcttgcc tcaaagaggt ccaggaagga caaggcggcc gaaagaacta 180
gttccgcgtcc ggagtacgac agtcaccgct ttatgagcgc tgtacaccag cagcgcttcg 240
aagccatcaa gggatggtcg tttctccggg agcgacgcgt ccagctcagg gacgacgagt 300
atac 304

<210> 140

| | | | | | | |
|------------|---------------------------|------------|------------|------------|-------------|-----|
| <211> | 376 | | | | | |
| <212> | DNA | | | | | |
| <213> | Glycine max | | | | | |
| | | | | | | |
| <400> | 140 | | | | | |
| | | | | | | |
| ctgaatgctc | tattcaatgg | agttgacaag | aataccttca | gactgatcaa | cacttgacac | 60 |
| gtggccaaag | atgcgtggga | gatcctgaaa | accactcatg | aaggaacctc | caaagtgaag | 120 |
| atgtccagat | tgcaactatt | ggccacaaaa | ttcgaaaatc | tgaagatgaa | ggaggaagaa | 180 |
| tgtattcatg | acttccacat | gaacattctt | gaaattgcca | atgcttgcac | tgccttggga | 240 |
| gagagaatga | cagatgaaaa | gctggtgaga | aagatcctca | catccttgcc | taagagattt | 300 |
| gacatgaaag | tcactgcaat | agaggaggcc | caagacattt | gcaacatgag | agtagatgaa | 360 |
| ctcattggtt | cccttc | | | | | 376 |
| | | | | | | |
| <210> | 141 | | | | | |
| <211> | 402 | | | | | |
| <212> | DNA | | | | | |
| <213> | Glycine max | | | | | |
| | | | | | | |
| <223> | unsure at all n locations | | | | | |
| <400> | 141 | | | | | |
| | | | | | | |
| agcttgctgt | agtataatag | gaagcaccaa | tataaatatc | ttgatgccta | ctattgatat | 60 |
| atagcattga | acatacatta | aactagctag | agagaataaa | aattgctgat | aatagtacac | 120 |
| tccataggtt | gtcatgatgc | gtaaaccact | actgcaagaa | aacactttat | attatagttat | 180 |
| tagctagcaa | tttttgtgtt | tggcttgatc | atgcttggcc | tcacatgact | gacaggtggc | 240 |
| gaatccatct | gcctatatag | acccttccnc | ataacttcct | ttnttactac | ttcttaagaa | 300 |
| aatttctaat | aggaaacagt | agaagaacat | taccatgaga | cttccatggc | tgagaatgaa | 360 |
| cggtcagttt | aagagacgta | ttgtgttaat | gtgttaaat | ga | | 402 |
| | | | | | | |
| <210> | 142 | | | | | |
| <211> | 391 | | | | | |
| <212> | DNA | | | | | |
| <213> | Glycine max | | | | | |
| | | | | | | |
| <400> | 142 | | | | | |
| | | | | | | |
| tgtaataacc | ttagatagaa | atgagagatt | atgttcttga | gttgaatccc | tcatgtgagg | 60 |
| tgaaaatctt | tagcatgtgt | cgtaaaacca | tagatatttt | ttttatgtaa | agtccaatag | 120 |

tgactagcaa aggtgaaatc cagtggcga cctggcttag tagaagattt aagtcttagta 180
aggaaatttgc aagggttgtga aacccaatgg ttgctggacc agttgcgaat tggttgtgtt 240
actgaaataa catctttaac ggtgaggatt ggacgcaccc caagggtgtg gtgaaccattt 300
atatagacct ttgtgcactt tcttctctgt ctctatattt tgctcttgc acaaattcaac 360
actacttttg tataaaaatac tacaattttgt t 391

<210> 143
<211> 378
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 143

ntggacaacc atatataaga agtactttca gattagtcat agttgacagc cactccggaa 60
gcatttcaag attgtcgtag tctgaaataa tcaaggtagcc taaggagttt gcagtttctt 120
gaagccattt aggttaaggcc accagctgtg gtaagccccca gaatgcaaca tatttttaacc 180
ttaacttgag gttatgctct tcattggcggtt ccttccacag atctaagtcc agactaacac 240
agtctttaac agacagagat tctaatttcag gaaaatttat aacatcctct gaccttgact 300
tcagactatg acaggcagca acattcaatg cttaagagc agggaaacttc acccctgcaa 360
agatagactc catattat 378

<210> 144
<211> 369
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 144

agctttcacac aaagaacata gtataggta aataaatata gctgaagatt taaatcacat 60
agaaaaacacc tttaatttct agtaaacata gctaattctc ccaactgctg ctcgctctgc 120
aattctatttca caataactcta catggaaaaa tgaggtatat tattttgtat ataaaatgac 180
aaatcaaact aatgaagaga aacgaagaan attagatacc cgagttAACAC actttcacaa 240
attaaacaat atctncttgt agacatatta catgagcata cctttgctac atctagnttt 300
ccaagctgta ttgcttagatc anatatgttag tcagggtcaa tagctacttc aagagcatct 360

tctatcata

369

<210> 145
<211> 397
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 145

tgtccacaaa natagttnt tgaagttgt catttcaatt tctcaactaa taaaatggat 60
cattttcaag gtccaatgcc ttanaatgat cacctcttaa agtaaaaaag aatcacttga 120
taagaaagaa ctacgttagt cttatttcct catcgcaatt gaggaatacg taggagcaaa 180
gggaaacacc cttgtcgact acaaaaagag aaaaatataa aaagggtgta aaggatataa 240
ggacttaaaa gggAACATAA aaaatcaagg tcatgtttgc acattcgatt aaAGCCTGCC 300
gtcccttggg acggacgtgt ggngtgctaa taccttcccc gtgcgtaaac acaactcccg 360
aacctttcac ttannagttc gtagatcgcg tcttttc 397

<210> 146
<211> 416
<212> DNA
<213> Glycine max

<400> 146

agcttcaata cgagtagcca cccatttact agatagttc ctaaccagtc aaaccattcc 60
caaaaatcatt ctgcagaata ataaaatgcaa aatagagttg gctaaacaaa aaagatcctc 120
ataatcattt cccatttgggt tcttggattc ccatacgact atacaatca ataattttt 180
aacagaaaagt atactcaa at tgaaacctag caatTTCTC aggttaaaaa aactacacct 240
ccttagcaatg gagaggattg ctcctacatc aaatgcttca gctggatcgg cctggattgc 300
acgttagctct tcattggttt cacgagcaac cttatacata agccagaaat tatttctaaa 360
gcattgcttt ccatcacagg caacaacaaa attaaacatt atacaaaacc gagaat 416

<210> 147
<211> 457
<212> DNA
<213> Glycine max

<223> unsure at all n locations
 <400> 147

ntganagagt tatctttga caacttctaa ctcccttcc tgtaattntt aacacgtgtg 60
 cccttacttt agtgtaagac ccattaaaaa ctatcaca aaaattgcac ctaacttcaa 120
 aattccacc gcctccactt agacctttaa gcttgaaac acaattccat aaaggttgg 180
 tgtcatcacc ttgttcttta acttgattag aagtactcat ctctacaatt aatcaatata 240
 ataaaataat aaataagaac acaatggcca attaaaata aaaaaattat gaatggtaa 300
 ctgttatatt taaaaactat tatcaaaata ggatataaaa taaaaatca taaagtatta 360
 aatgtattag taggtaaagaa agtaaagaan aataatatta aactaaaaat tcttatgagc 420
 ctacgaaaga agaanaaaaa attataaaaa ttggaaa 457

<210> 148
 <211> 396
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 148

agtttcatgc ttaactatgt atggaaaaac ttcattactg ttgttcaaga catacaagtg 60
 agcttgcac aaatcttcta cacttggagt gatcacctgc agtcctcttgc aacccttacc 120
 acccactctg tcacatcgcc gacactcagg aagcccaaca gcttagcct tctctaagta 180
 ttctgaacaa aattcaatgg cttcttctgc aatgtacctc tcaacaatag atgcttccgg 240
 acgatataga ttctttgtat accctttaa gattttcatt tatcgctcaa ccgggtacat 300
 ccaccgtaga taaacaggac cacagcattt gatttctctg accagatgca caatcaagtg 360
 aatcatgatg tcaaagaaaag cangggaaaa tacatc 396

<210> 149
 <211> 462
 <212> DNA
 <213> Glycine max

<400> 149

tcaccaccaa gacagtgtct tggataagaa gcttagagag gatgcttcaa tagaggaaga 60
 gaatgagaga gaaagaggaa agggcgtggg aattgatgaa ggagattagg gagagaagtt 120

gaactttgaa gtgtgtctca caagtttctc attcatcaaa gttatgagaa gtgttacaca 180
tgtttctatt tatagcctag cacaatggaa gcttccttgg gaagctaggg gaagaaaagct 240
tccttgagaa gctagagggg gctactcaca cctctccaat agctaagctc accccatgtc 300
aagatgcattg aaaatacaat gggaaacttc cttagaggt aaggtagctt ccttaggaag 360
caaggaagaa agtttccttt agaagataga gaggggctac tgatgcaatc ctaccccaca 420
agggcattgg atagaagaat ccaagttagat tggctagag at 462

<210> 150
<211> 406
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 150

agctttttt tacgggttctt attaaatccc atccccttcc ataagctaaa atactcttc 60
atatacagtg atatggtcag acacaaaaaa aagtggaaaa tattcaagtg taacacacat 120
tgtatttgaa aagcaacaac tagtttaaa gtacaaaaca agcaaaacta aaaattacta 180
cttgctagta gttgaaactc cctagtaatt agcattaaca ttggatttat tcaaattcac 240
ctaaatacaa catactaatt aacacgaggg tagcgttata tctcaaacat taaaagtggc 300
ataaaaacaaa ttagcaagac tatnttagtt caactgctt ctc当地acta actctaatgc 360
attctcactg cttcgtaa taaacttggg tcaatgctag atcact 406

<210> 151
<211> 440
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 151

tatcttggac ctc当地actgaa tgttagttgaa cctcagcttgc ctgaactcac ttcccaaccc 60
atagctgaag atgccttgc aacaatttgc acttccagta ttccatataa aatatctcct 120
ttaaccacct ttgcttcaac ggggtgtct aaagaaaagga ttcaagaaat gttgtgcctt 180
caacaggcca acctagaaga aaccatggaa aagagttcc tcttttaatg catggaaaca 240
aatttgctta tcaagaacac cttgaaagct tcactcaagt cttcaacga cttgttttg 300

ttcaatatga atatgctaat gcaacaacga ctgcctccaa ccgtccaacc ccttctgcca 360
ccaccagcta cttcttccac actcacaact tctaaaanaa ttgtgctgtg cctcgacctt 420
caccaccta gtcaccacca 440

<210> 152
<211> 395
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 152

agtttgtaact ctgttggag gaacccttac tcaagctatc ctAAAATGAA tacaacaaac 60
cataAAAATT tcacacact catagcaaAT cagaAAATAAA tggcgtGAAT ggTTTATCAT 120
tacAGTTACA acAAATCAAAC cacaAGTGGT gcacAAATCA aacATGAATG gtgcGAATGG 180
ttttCTTG caactactat ccacanAGTg agatAGAGGT tagggattAA acAAATCANA 240
gcacacaAGT AATAATGTTA tttgtAGTGT gagAAAATAA caATTGAAGA ATGCTAAGGC 300
gcgaAAAATAA ttcaattaca cttctccann AAATGAGATA aagCTTCAGG ATTAAGAAA 360
tacaAGTGNAA AATTAAAAA GTATAATGGT ATCTG 395

<210> 153
<211> 376
<212> DNA
<213> Glycine max

<400> 153

atacgttgct cattgactct cgattgtac acagaatgac caagatctt acggatct 60
gcagaAGAGC atagaccaca gactctgCG acatgtgtAG atttcttatt catggaaAGA 120
cgatgtacta cggTGaccAA cggatGAAGG tctcTTcAG gcttttatt ttcaTTGAC 180
gaggAGGAAT gtgcggccAC ctgatggACT CCTCTAAGAA caatGACATA atttcttGCA 240
ctgaattGTT gagAGTTGGA agccatCTTC tcaatCAAAT tcctAGCTTC agcaggGGTC 300
atATCACCAA gggCTTCACC ACTGGCAGCA tcaatCATAc tcctCTCCAT gttgctaagg 360
tcctcataga AATATT 376

<210> 154
<211> 264

<212> DNA
<213> Glycine max

<400> 154

agcttagaag aattaaaaat gaaaaaaaaa actataataa ctaaaattgg aaagacgtcc 60
acttataagg actaaaatta gaaaaataaa ctatagaga ttaaaaatta aaaaaaatgc 120
taacttacag ggacaaatac atatthaagc ttaaaaataa cattattcta aaattaaaat 180
ttgggctcct agtagcatc aaaacagtcc atttattaca attaagatca agccagagat 240
acttaaataa aataaataaa aaat 264

<210> 155
<211> 314
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 155

tgttacaaat gtggtaaacc ttatagaaac taaaggctgn tcgagtcgtg tggttgcc 60
ggataaacat ggcttgcag cagccaacaa aggctcgcaa tgacgatgga gaccagtgtg 120
atgctcgctg gagatgcgtc acgcgatgtg tgcgtgaacc tcacacgcca aggccctgca 180
tggggttgcg tgcggcggtg ttatgagttg gctgcgcaca tggagtcgtg gacgatgtgg 240
tggccttgg cgatgatgat gggcattgtt tacctgtgaa aataaaaagt ggcaaggctc 300
accacggacg cctt 314

<210> 156
<211> 403
<212> DNA
<213> Glycine max

<400> 156

agcttaacaa tcctttgat ctatccaat atattctat ccctatcata taacttgcc 60
cactcatatc cttcattta aagttacaag agagaaactt tttcggtgag cattttctta 120
aattggaaat tgtgatgttg agcattttc catctaaat ctctctagta ctttattgat 180
atatgcttt tgagacaagg ttaacaatcc tatgatctat ttcagaatat ttctatccct 240
atcacaaaaac ttgcctcacc catatccttc atttcaaagt tattagagag aaacttctta 300

gtctcatgaa gaagatcaag atcattagtt gcaagcaata tatcatcaat atacaggatt 360
agaaaaataa ccttactcct actgaccttc agatatacac att 403

<210> 157
<211> 450
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 157

gtatccaaag ctggacttat atacagtatt cttggctcta ccatatacta tggctctaac 60
ctatcgacga acttattgga agaatccctc ccgacttagca tatatgatcg tcttaccac 120
caacttctct ctaagctcta ttagtcgtta ctctcttgat ttacggatg ttgtgagcgc 180
cttggcttt gacttgagta gaagtactca tctctacaat taatcgatat aataaaaatac 240
tcaataagaa cacgatggc aattttaaat aaaaaaatta tgaatggtta actgttatac 300
ttaacaacta ttatcaacat aggatataag attatgagtc ataaagtatt agatgtataa 360
gtgggtaaga aagtaaagaa tgataatatt aaactaaaaa ttcttatgag cctacgatag 420
aagaangaaa attataaaaa tcggaaagta 450

<210> 158
<211> 401
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 158

agcttccaaag aatcaagatc aagattcaag actcaagatt caagaatcaa gagaagactt 60
aatcaagata agtatgaaaa ggtttttaa aaaattgagt agcacatgga ttttctcac 120
aacatgttta tcaaagagtt tttactctt ggtaatcgat tactagattt ttctaatcga 180
ttaccagtag caaaatgttt ttgaaaaagt tttcaactga atttacaacg ttccaattga 240
tttcaaaaaag ctcttatatg tttggtaat cgattaccac tgtcttgaa cggtgaaatt 300
caaattcaaa tgtgaagagt cacatccctt cgcataanag ctttgttaa ttgattacac 360
tgatttggta atcgattacc agtgattgtt tctgaataaa t 401

<210> 159

<211> 387
<212> DNA
<213> Glycine max

<400> 159

tcttatccaa ggctcatctt ggaggcgaag ctacttcttc catggcttat tccctaattg 60
aaggcgecta ctctcagctc ttctactttg tcttccgctg catctacatg gtggaaaatc 120
actattaaag gacctcattt aagctcacag atccaacctg catagagacc ccacaggcaa 180
gcttccatca taaccactct atttgccta ccagggatat ccaacttgga cactgcactc 240
gccaagtaca tacacgacat acatcattac aatgacacta tcaacatcca cagcatctaa 300
gtctgatgac actatgatca tctacctgat cccgtctcgta tgtcattctc aacatcaaca 360
gtatctgatc tcaatgacat aatcaac 387

<210> 160
<211> 411
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 160

agctttccat gaatcaacaa aatgaataga taactcacaa cactattggc atccctgcct 60
ccacaaagca gaagaagtcc atcagagcgt gcacttgcag ttgcataacct agtcaaagat 120
taaatggtaa aatgaaaatc gcaccatgac aagttgcacc aaatgatagc caactaatat 180
attnaaggta agtctacact tcacagaatc gattgtttac atcttctaatt gtaagatatt 240
taaagtattc tcttgctt acaaaaaaca aaaagcaata atgtgtaac tttgttccac 300
aaacagtaat attaatgcta atcaaaatgg ccaaataaga tgagattcaa ttcanaccct 360
actcaagctg cagtcttaact caagttntcg tacagaanat canaagaaaa t 411

<210> 161
<211> 442
<212> DNA
<213> Glycine max

<400> 161

tgttagactga atctataccaa taaaaaagat ttgtgttcag tacaccgctt atcataattt 60
tttttttctc tagccattat ttttagggag gtagtttagc taactcacgg attttaattc 120

tttacttgca tgatgattca ttttccttt ctatacacat tgttttttt gaatgatttt 180
tacatatgtt ttgaatcaaa catttcagaa ttatcattta tattaacgct atttatccgc 240
atagttaat tggccatcgt caaactaaa tatacggcag atatatatta taacttttg 300
tataatacat gatTTTtaac aaaatctta ttacatTTT cttatgataa gggatttagaa 360
ctttttttt gtggacaaga gatacaagtc ttcattcct ggaacatata taaagctgaa 420
tgaatatgaa atgcctccc gc 442

<210> 162
<211> 324
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 162

tcaagcttca agttagaagt tgaacacttt tatgttagaa gtgtcaaATG aggAAATCTG 60
aagcgTTGTG ttttctatgg ggaactacaa ggCGCTAGGA gtagatgggt tccaatctat 120
tctctataag agcaaataag atgttggTGG tgaAACCTT tGTACTCTTA ttAgAAGTAT 180
nttttataat cccactaaca ttgatgattt taattatact cttattgctc ttaattccaa 240
gcaagatgtg ctcactaaca tgaAGCactt taAGCGcata atgttgca atggTTccta 300
cagacaatga caagaatATG tctg 324

<210> 163
<211> 379
<212> DNA
<213> Glycine max

<400> 163

ctaaacgatg ggctagctt agcttagccag gcaactttca tggTCTTCat tagagAAact 60
agctcagaag tggTCCCTA atgatctAGC ttaAGCTAGC ttggCCACTT gcaaATTGtG 120
tacactgttc' tttcaatgtat agcttAAAT atctcttcaa agagatcctt actgttagttc 180
ctacAAAGAG actgaACGAC atAAACCACC tcacAGCTAG tacactaggc tcttAAATA 240
tttctctaaa gctgagtttA ttcaAAAGATC AACCCAAATTG tgctcaaaca atgttcagaa 300
gcatgagAAA catatcatAG ttgtcacAAA aatcgcaAAA aacaAGTAAA agaggtaatt 360

ataattgata tcttaactct

379

<210> 164
<211> 173
<212> DNA
<213> Glycine max

<400> 164

tgcttcata ctttatacaa gaatgaagct ctgataaccac ttgttagaca agtggcctca 60
catatattaa gaaggggggg gggggtccca tcctagacac ccgcttcatt ctctcctcat 120
ccttatccaa aagtattttat tctattctac gcctccata tattgatctt ccc 173

<210> 165
<211> 377
<212> DNA
<213> Glycine max

<400> 165

tgaaggatgt aagattctgt gattctcaa tgtccaccac aatgtgatca aactttgatg 60
tcagtgttct gagcacccctc tcaatcacca gttgttcatt aatttgcatt ccacagcact 120
tcatctaatt ggtgagtgtg agaatcttgg taaaatactc agctactgat tcagtctcct 180
ccattgcaag aagctcatac tgtcttctca atgtctgaag cttaaccttc tttatcttt 240
ctccaaagat aatccaatat ttattnatca caagatatact aaatataatat tttagtaaag 300
taaaagatag atacaattgc tttaatatata tatattgtt ttcttattct cgaaagatgt 360
tattatcata atatatt 377

<210> 166
<211> 449
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 166

tgttcctana tgatggctaa gcttaagctaa gcctggtaac tttaagttc ttcattagaa 60
tagctagctt aaaagtctgc ccctaatacgat cttagcttaag cttagcttgtt aacttccaaa 120
ttctttacac ttttcttca atgatagctn taaatatactc ttcaaagaga tccttaatgt 180
aattcctaca aagagactaa acaacaaaaa ccacaaaaaa gcaataaaac taagttctta 240

aaatatttct ctaaagctga gtttattcaa ggatcaacca aatttgctc aaacaagg 300
cagaagcatg agaaaacatat aatagttgca aaaaaaattg caaaaaacaa gtaaaagtgg 360
taattataat tgattattta actctagtaa aaaaaaaaaag cattgatcgt ctaaccttat 420
tttatcaatg gttaaatact taattcaat 449

<210> 167
<211> 331
<212> DNA
<213> Glycine max

<400> 167

agcttggct tgatttttt ctaagttctt taacaagatt agaacaatat acttgcctt 60
catthaactg tcttggct tggcgccac gatcaacaaa gtacttcga cacctactat 120
atgttgattt gaccaacact gttatcggtt tggcgaca atccttcaaa accttattta 180
tacattttga gaggttgggtt gtcatgtggc catatcgacg tccttctcta tcataagcca 240
tcgtccaatt ttcccttgaa atacgatcaa tccatgtgc tatggctgga ctcaagttgaa 300
cggaattttc taaattttga ttaaaaaaaaa t 331

<210> 168
<211> 449
<212> DNA
<213> Glycine max

<400> 168

tgtgaagtgc cacgttagag aacagaacac tctttcttc tctctgttg aacaggctct 60
tcttcacgag aaatttattac tctctctact ttatcattag ctcttatttc agtgtttgac 120
ttttgcgtt gttgttcgt tgagtgtgca ctcggttctg tttggcgtct ggacaaaaaaaa 180
catgggggaa gaagaagaag aagaagttag cgtcaccgct ttgcacatc caaacagcgg 240
aaacgatgat cagagcctcg aattcgat atatccttg agcagttact atttggatc 300
caaagatgct gttccctcca gagacctcac cttagatgat cgtgttctca ggatgaagta 360
caagctcggg ttcttcttc tcctctctgc ttatctac tgcaatataa acactcttt 420
tgttctcttc tttaactatt ctttcttc 449

<210> 169
<211> 394
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 169

agcttggtctt attcatcttt ntcattctct tctccctttg ccaacctgaa ttcttttgtg 60
tctcccttctt ccctttccaa gagaattcaa aggaccccgc ctgagaattc ttttgattct 120
ttcctttccc ttaaacaaaa gatttcaaag gactaactgc ctgagatatc ttttgttcc 180
ccttacaaag attcaaagga ctaaccgcct gagaattctt tgtcttaaca cattggaggg 240
tacatccctttt gtggtaacaag tagaggatac gtctacttgg gttgttgaac taagaataag 300
agagggtaca tctcttggg atcagttcaa gtggagggtta catccacttg gttgttcaaa 360
gagaacaagg gaaggtacat cccttggta tctt 394

<210> 170
<211> 463
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 170

tgtgcttaa gcccacttaa caaaaaactg tttttataaa gtaaaaaaagc cacactctt 60
acaagttcta catatgttgg caataaggc caacatatacg ctctctcatt tacagtaaac 120
aaaactttaa aatttggttt aggcctcaact cacccttggg ttggatcaat tgccaccatat 180
aaatatgtgt tccccgtgga ctaacaaatt ttatggct tttatcggtt catttagacc 240
tttgcatgat ggcgactttg atgtcataca cattacttgc gctttttt cttacacat 300
ttgttgcatg tcttcaatct aatttggta atctggaat caggttgaa gttatgattt 360
caaggaaaac agatttgcgc atctacacccg atctgcgatc agctntccag aatcaagggtt 420
cttctatgcc ggcactcctg ctacatcaaa tgcanaaagca gct 463

<210> 171
<211> 417
<212> DNA
<213> Glycine max

<223> unsure at all n locations

<400> 171

agcttgcttg tggtgcttct atggaggcgt gatcttttag cttcaatgag gtccttaat 60
ggtgatttgc tatcatggag atgcagcaaa agacaaagga gaaaagggtga gaagaggcgc 120
catccaatag gtaataagcc gtggaagaag gagttcacc accaagatga gccttgata 180
agaagcttgg aaggatgctt caatggagga aaagaaagag ggagtgaaag agagatgggg 240
gagcacgaaa ttgaaggaat aaaagaggga gagaagttga actttgagtt gtgtctcaca 300
agactctcat tcatcanagt tacaataagt gttacacatg tttctattta tagactangt 360
agtttccttg agaagctctc ttgagaaaac ttcccttgaga agtttctatg agaaaaac 417

<210> 172

<211> 461

<212> DNA

<213> Glycine max

<223> unsure at all n locations
<400> 172

tganatgagg aagtgtggaa gggtgagact tcctactttt attcggtggc cacagagtgg 60
tacctggaga tatgtcgccg nggtcaggag accttggggc cgtcaggtgg ggtgctattg 120
ccccaaacca agcttgcacca atcccgcaccc aacccgggca tagtcagtca gtgagaacct 180
gtgatgtacc taaacaggcg agctccttggc agtcaatcga taaaagaaca aagaccacaa 240
agcaaggagg ctgtgtggc ggctggccag ctgtgaatct tgagtgatat atgggatagg 300
gcctctggta atcgattacc gaggggtgggt agtcgattac aaggcttana agtgaagaca 360
ggaagctaag atggcctctg gtaatcaatt accaagagag tgtaatcgat ttccaggcctt 420
annaacgaga tcaggaagct aagaggcgtt ctggtaatcg a 461

<210> 173

<211> 260

<212> DNA

<213> Glycine max

<223> unsure at all n locations
<400> 173

tttcttctat gattcattct atgcaccat atatcggt gttggatggc caacaattgc 60
attctcggtg gggataactaa tcactgcgtt ttgcaataact ctcactccat agtaacaaca 120

caagttcaa agcaaactt tggtgacaca agatgataca gtgcgngcta gttgtatat 180
tcattagctt aggataaaat caattccgac cgttcggtcg tgccgtaacc acgttggaaa 240
tcaaagagag gtaaaaatg 260

<210> 174
<211> 237
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 174

tagtagaaga tanacactgc tagggacgaa cctggcaatg aagacatgct gtctactcta 60
tatcggaacc ggtgataggt tcaaacctaa ctatttatt gatgataata tctagcagga 120
cctatggcta ccatggaagg atgcagtgtat tgtgacacta ttatgaatac tgatatggat 180
tgtcacaatg cacgataggt tgtaatctat ttggacatta tcacgaagag gaattaa 237

<210> 175
<211> 324
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 175

agcttggaaat tgaacaacgg aagctctcgaa gaaaatcgag tggtcataaaa ttttcacaca 60
gatgtccgat tcggggaaat aatatatcgaa gacgcacgaa attgaacaac ggaagctctc 120
gagaaaatttgc aatggtcata acatttcact cgatgttcg atccggggac ataatttatac 180
gagacgctcg aaattgaaca accgaagctc tcgacanatt agaatggtcg taactttca 240
cgcgaatgtt cgattcgngg acataactca tctagacgct cggaaattgaa caacggaaagc 300
tctcgagaaa tttgaatggt cata 324

<210> 176
<211> 381
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 176

ngccgccacg gagttntccg actatgctct tgggtgttgg aacaagctac aaaaggagag 60

agcaagaaaat gaagagccaa tggttgatac atggacggag ataaaaaaga tcatgaggaa 120
gcggtatgtt ccggctagtt actcaaggga cttgaaattc aagctccaaa aactaaccct 180
aggcaacaag gggggttgagg agtattcaa ggaaatggat gtgctcatga ttcaagcaaa 240
tattgaagaa gatgaggagg taactatggc tcgatttctt aatggttga ctaatgatat 300
ccgcgatatt gttgagctgc aggagttgt tgaaatggat gatttgcttc acaaagcaat 360
ccaagtggag caacaattaa a 381

<210> 177
<211> 276
<212> DNA
<213> Glycine max

<400> 177

agcttattgg taaaaaggc gttatttcaa tcacttagat aaaggtggaa gttataactt 60
cctccattat taattaatct tcttcaccc actctctcca tatataaacc caccctcaat 120
gctttcaaaa agtttaaaaa ataattataa gaaaaatggc aacatgtaaa ctcccaccta 180
cctaagttca gacaaaaaac acccacacac acaaagaagt taaggaacaa attggAACCA 240
taagcattct ctcttagtgg agaaggagaa aataat 276

<210> 178
<211> 459
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 178

tannagttga aaataatata aaagtcctaa tacaggtcca gcatgattcc actcactagt 60
acaaaaactca cctactaaaaa tgtgtgctaa ccagtgtatc atacagtggc ctccattcaa 120
tagtttagtga aatctttta ctatattgt tgagtagtct aactaataaa tttccccatc 180
taacctaata aacaaaaaat ggcacaatca tttaagagca agcagagaca tatagagaac 240
aagaaaaactc agcatctcat ctaagacaaa atgcagcagc acctaagcat aatgcttatt 300
acgggacaca tgagatatct caaggccctt ctgaataagg ttgtgaacat cctccanagc 360
tacatcactt tttccttta tgaaattgca caacacacat aaaaagtttag attcaaccc 420

aacaaccaac tccatctgga tacaagccat gagtgaaaa

459

<210> 179

<211> 330

<212> DNA

<213> Glycine max

<400> 179

agcttgtaat cgattaaact gatatgagac atttgtctgc aagcttcaaa cacttgtgt 60

actggttact atcagtctgt aatcgattaa aacagaagag atgtaactat agaggaaatc 120

ttctaacttt agaacttttc ttctaactcc aacatgatga tgcatgatac acatatgaaa 180

tgatagagac aaagatgcaa cacacagtac aataatcaat acaaatgtca ttcaagagag 240

ttgggcatgt agaagacaat aagatcaagc tcttctttaa gctgtaatgc taagtataca 300

tggtgcttcc cctatctcta acatgcaata 330

<210> 180

<211> 457

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 180

taatcttaca aagctctntg aaagaagggtt ttctatgtat tgattccttc aagtatccat 60

cttgagatac ttttcttgaa gaggaggaat atgcaattgg agtgtatgga gaaaggaaac 120

tagtgctcaa gcaatttccc atcagactca ttattgctcc taacccaaac agcctgatat 180

ttaagtatag gccaaactgca atgaagggtt atcccccttgg tcccaccatt tttgtctttg 240

tgggataccc taacggaact aagggatact atttctacaa cacatatgag gaaaaagtat 300

ctatcactaa gattggagtg ttcccttgaga agcaatacat ctccccatgga gccaatggag 360

gaatgttagat ctggagaaa tccaagtggc acataaacatt gatccaccta caatgaaaca 420

agaattgatg ccacaaaagc attgttagatc catcttt 457

<210> 181

<211> 368

<212> DNA

<213> Glycine max

<400> 181

agctgtggaa agtgttggc tcacccatctc gctaaggccac tctgatggct tatcgacgt 60
ctgctatatg caacattcct gggctaagcg caaggaagaa tccataagaa gatgagctgt 120
acaagtgcgc taagtgcacg cgcttcatct tactaagcgc accacttgag ttcatctgct 180
aagcgagaaa agcgggctaa gccaaaaatc actaacgtgc gctaaggcggt ccataagtgc 240
gctaaggcaca cgagcacaaa caaggccgcc tagtttaagcc tgaaatcaga tcttgtgaaa 300
ggagtatgga ctaagattca gagctctgca tgccttagggt ttcttagagag agaaagtgc 360
agttctag 368

<210> 182
<211> 377
<212> DNA
<213> Glycine max

<400> 182

gctattacgg acctataata ctcagctgtt tatccgtttt tgtgcaagac atatttaaac 60
cgatcaatttgc tcatttaagg cgttggacca ttaacgatct ctgggttttt taaaagtagt 120
ggtaaaggta gacgtttatt gtatgttcc gaaggtgcattt attaaccat aaaaggcagag 180
agaacctttt aaggcattgg accttaaaac ggtttttagt gactttgcg gacaaaagct 240
tcatttgaga gttgatttttgc gccttaagttt cactttgggtt attagtcaat tcattcaagg 300
aaacttgcaa agaaaaatgc ccgactgagg tttttctttt ttgagattgtt attcaaagat 360
attgcgatttttattt 377

<210> 183
<211> 364
<212> DNA
<213> Glycine max

<400> 183

agcttagaat ggccgaaagg gacgagtcaa gggtggtaag catggcatta aaggataaat 60
tgaaggcttg tcataaggtaa aagagaagtt tgaccgaaca atcgagtgga acggaagaga 120
atatgttgac gatcattgtat cagtataagg agaaggtaaa cctagctgct agtcttaggc 180
agagactaga ggatgatcat gcgaaggat tgactctaca aatggaaagg gaagcaagag 240
agagggtgat agaatttattt cgtggaaatg gatggataga ttcgctctca 300

ctctgaatgg gagtcaagaa gcttcagggt attagccaga accaggcaat gacggaagt 360
tact 364

<210> 184
<211> 439
<212> DNA
<213> Glycine max

<400> 184

gtctccacta agttgcctaa tgccctgaaat gtctttctg atggcagagg tccttagatgc 60
agggaaagaaat ttctccaaga acacccttta aaggtcatcc cagttaaaaa tggacctggg 120
agcaaggtag tatagccaat ctttgccac tccctccaga gaatgaggaa aagcctttag 180
aaagatata tcttcttgc catcacgggg cttcatggtg taacaaacaa tatggaactc 240
cttaagatgc ttataaggat cttcacctgc aagaccatga aacttgc acaaatttat 300
tagtccagcc ttgagaacat aaggaacacc ttcatcgagga tattgaatgc acaagcttc 360
ataagtgaaa tcaggtgcag ccatctccct aagagtcc tcacgaagtg gaggttagagc 420
catgttctca gtatgaaaa 439

<210> 185
<211> 396
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 185

agcttggaga ggatgcttca atggaggaaa agaaagaggg agagaaagag ggaggggg 60
gcacgaaatt gaaggaagaa aaagggagag aagttgaact ttgagttgtg tctcacaaga 120
ctctcattca tcaaagttac aacaagtgtt acacatgctt ctatttata tag actacgtac 180
ttccttgaga agctntctt agaaaaactct cttgagaagc ttctttgaga aaacttcctt 240
gagaagctag agcttagcta cacacaaccc tctcataact aagctcacct tcttgagaag 300
cttccttaag aagattccta aagaactaaa gcttaactac acataccct ctaatagcta 360
agctcacctc ctgagatgag aactagagct tactcc 396

<210> 186

<211> 139
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 186

tcataagcat ctacccattg cgtcaanaga tacatgcaaa gtgctacaca tgctgcttt 60

tatacactga ctacctacct tcatggcctt tctatgagga cactaccatg gaaaacccta 120

ttgttaatatc taccttgat 139

<210> 187
<211> 396
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 187

agcttcaaag aagactattg aagtgtggtt caatcaattt acaaactttt ggaggcagta 60

gacaaagatg acagctggga aaggacaact agcagagtca ttctgaggtt gtcatgttga 120

ggaaccttgc tcatgagcat tttaaagtaa aatcagaatg gtgatgttag ctgagaaagg 180

atccataactt gaagtagatc cttttgatga agatcaacta ttatgtctt ctgaagtagc 240

agaactgcat caattntaat acatcttcac aacacanagc atcctgaagt agattgcttc 300

attaaatcac agtgaaggca tagnttcttg gtgttagtg gtctatcana agttagaatg 360

tgttaaagtct tcaatcttcc attctgatga tccaca 396

<210> 188
<211> 395
<212> DNA
<213> Glycine max

<400> 188

tgacactatc caagactcta tacaatactg aagctctggc ctctacagat ctccacacag 60

cacaagtact cttaactctc tggagcttgtt acctttctct ctctagaaac ccttagacatg 120

cacagatatg aattctaatac catactgccccc ttgtaaaatc tgaatatacg ctcatatatg 180

cggccttggc cctgctcggt cgctgtacgc acttatggac cggttaacgc acatttagaga 240

atttacgatt acagcgtgcc tttctcgcat atcgaatgaa ctgaatacgc gcacttaacg 300

agatgaagtg gtgcggtcac agaacgctta ccaatcaact ttttccagag tcttgctcgc 360

acttaaccctt tgaatgttgc gcttatcgga cactt 395

<210> 189

<211> 499

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 189

cgattgaccc tttgagcccg agatcccggt agtcacctgc agcatgcgag ctatgcagca 60

attgaaatgg tcacaactgt tctctctgtt ttacgagagg ggcacataag atattgagat 120

gctcgaaatt catctatgga tactcttgag caatacaaatt ggtcagtgac ttttcactgg 180

gaggtgcgat acacgctcat atgatatcgc gatgctatac attgaacaac agaagatctc 240

gacagattca gaaggtcata tcctntcaact cagaggtctt agtcaggccc cttagcatatc 300

gagacacgaa tattgagaga acgaatggtc tcgacaaatt catatggaga gagatttca 360

cttgtatgtc tcataatgc gcataaggaa taaaaacgct cgggtctgtt gatggagagg 420

tctctagaaa acaaaggggc gttgtcttgc acgcatggca attcagcaca gagtgatgtg 480

acctcgatata tactcttcn 499

<210> 190

<211> 428

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 190

gaggaagcaa ccctgctcgc ctggcgagc tgagctgcc tggcgagct gggcgcaac 60

cacctccccct atttgctat aaatagggga ggaaatgaag aaggaagggg tccggccccct 120

ttggcacttc tctcttttc gaatttgctt gaaaaattt gttccgtgaa gaaaatctaa 180

gccgaggcgc ttccgaaacg tttccgtAAC gtttgcgtg aggaatctag cagaggtttc 240

aaccgttctt cgacgatctt cattcgataa gcatcgatct tcgaccttct gcgggtaagt 300

accgccaacc aagctgttca atgagagttt tggacctcggt tggcttcaca ttgcactcg 360

tgctatattt tcctatagcg ctgacactag aacangaagc tgatttggtagcaagtaa 420

ataagact

428

<210> 191
<211> 238
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 191

aacattcttt tcagctaang attatatact tatgttcttt tcaaatgcan aatcaactaga 60
tggatgaaca ttagtgcatt gacatcttgg cttctagcac aacaggagaa atgttaacgc 120
ttttctccgt agggcatana aatcaatcta ttgcatggga tcactttga taaatgccat 180
accccaatcc tttatccaa gctaaatgtc gacattgtgg tgctntgatc aaatata 238

<210> 192
<211> 210
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 192

tcatatatcc canaacccca tacncacgan nattaagaga gaaagaagtc cacccagacc 60
tggattntcg aagtcccact cgtagccacg cacttcacga ccccgaanat gccctccttt 120
cgcgatttgg agcagaaaatg agcacccaaag gtggagctn ntgtcggggtt tcaatggaga 180
atggaggaga agaaaaaagc aacgtgagga 210

<210> 193
<211> 220
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 193

gagtttcgaa acaagagtat aaagtgtgca gagtcaaatc tggattttgg ctggatgan 60
natgggctag aatatatcta anatcagata cggnnnggtgg actgcacatcg gnntcgaaat 120
gaggganana aatatgccta actcaactnn taaaaanaat caactaaggc taccggagta 180
ctatacgaac tntggtcata tgctntgatg atgagagatc 220

<210> 194
<211> 209
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 194

attattcaga tgcttatga atgacaatca aaggtgactt ggaaacaaga aattaaagag 60
acttttcatt tgccaaacag ntntatcctc tcaaaaagaa taagaagtnt tctgaactga 120
aatggtnat cctctcanaa agattcttg gtcaaccact tgcattatca ataanggaat 180
ttgattgatc ttcatgtac aatctatct 209

<210> 195
<211> 194
<212> DNA
<213> Glycine max

<400> 195

actactcata attaatgcta ctagaagagg atgactgata actattatta ctataatatc 60
caacaagcta ctcaccattt aggaagttga tctgtattac ctgaacaatg gcaactatga 120
caatgatatg ttatgttcaa acattatacc ttgagttagg ctatgctgta gaaatgagac 180
taattaatgt catt 194

<210> 196
<211> 366
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 196

accatggtaa tcagaataga aagtctactt cttatgtaaag agttgcagag attccttct 60
cacttctcac ttattacagc tagaacctat gtctagctga tggttctatc ggctgtgatg 120
caaagattca ctatcaccct gtatctaata taaaagtggc cagattcaga attgttactt 180
tagaccacgt aaagaataat atgcataatg gacaatatac accaagagaa actcagatag 240
cacaatccag tggtagagaa gcaacactta atttagtggt canagtcgt atatagacat 300
gagaatagat accttgccan aattatccat aagacatttta tttccattcc ctaccagtgt 360
attcat 366

<210> 197
<211> 276
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 197

atgccctctc tctttctcn tttctcttc tcttctggct ctcctatatt cgnttagn 60
tctaggctct tcttagacac tnnttcatt ntgcaattcc actnttagta ataaaaattc 120
gctcttcaat ctataatttc gttcttatt gattaatgca aggctaagtc tccagcgtct 180
gtttctcttg aggatcaagc acagttctct ctgaggtctt attattactg gtaaaattct 240
gntcagttt tcttttact acataactctg aatttg 276

<210> 198
<211> 234
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 198

caataatcaa taatctatct ttcaatcttc tctcaacatc attcaatatac tntcaactct 60
ttctacacna atttctgatt catttcttta catcttctta aaagtttttg ttcaacactc 120
tctcttatga gaaaagttct ttgttcanaa acttgtgtta ttcatcctt tcattcttt 180
ctccctttgc caaaagaatg aaggactaa ccgcctgaat tctttgtgtc tctc 234

<210> 199
<211> 499
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 199

atgaccctta gancttcaga acnctctaga gtccacctga ngtatgcgag ctccagacag 60
tttagcaatcc tgtggtgcat tgtttatct gaactaccat tatcaagtaa tattggAAC 120
accatgccat taattgtacc actaaacttc atagttccta ggccagagga accttaagg 180
gcattataag atagatgatg gtctaaattc atatgtgcca tggcttcctc aataatagaa 240

ggtgtgctcg gttgcacttc agcatttca tcctcatcca ttgcaatagt aggcatggc 300
gattatgaca cctatgataa gaagtgaact tctcatgaca tgtataacat agccctttct 360
cccttcgaat ctgcatttca gctggtgaca ttcttctaac atttcctggg atcggaaaac 420
aggagttga tctaactcag tctaggcacc tgagtccaca ttcccttgaa tggatatgct 480
aagagagaga cttgtgact 499

<210> 200
<211> 299
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 200

tgtcgctaga gctgacccat caactgtcct atctctntca gactggcgac tcttangctc 60
ttgatcttga cttgatagaa cctctntnta agtgaagggt gcctgactcg atcccattgtt 120
tactaaagtg gaataaaaaac cagtgcgaat caagactgtg acatctatca caggtgaaat 180
ggatgaatgc ataaagaaat gcatatggca cagatgccat ttacggatac ganagcccga 240
gagaatatct atttcttana tacaacattc nggcagcata gtgcccgtatg catgcattt 299

<210> 201
<211> 429
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 201

attattatct gaactaccat tatcaagtaa tatttgaacc accatgccat taattgtacc 60
actaaaccttc atagttccta ggccagagga acctttaagg gcattataag atagatgtatg 120
gtctaaattc aaatttccca tggcttcctc aattctagaa ggtggttctg gttgcacttc 180
agcattttca tcctcatcca ttgcaatagt aggtattgcc gattaggaca cctatgagaa 240
gaagtgaact tctcatcaca tgtataacat agccctttct cccttcgaat ctgcatttca 300
gctggtgaca ttcttctaac atttcctgggt ntggaaaat tgganngtgt aattcatttg 360
gggcttgaa gtaacggcgg catgcttgtt gactttctt actgacacta ccgctgactt 420
ctacttcct 429

<210> 202
<211> 433
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 202

atgactattg aataatctat tcatgttcc tttgatgaat ctaatgctat ttctccaaga 60
aaggatattt tagatgatgt tgcagaatct tttagaacaaa tgcataattca tggacaagat 120
tctaaaggga aaggaaagg aagcaatgaa gatcctccag aagaagccaa atcaaatgt 180
gaagggtggca tccatggagt tcactgacta tgcttttatt tggtgggact aacaacaaga 240
agatttggag aaccttgggt gaatacttag gaggacatga aaagattaat gagaagaaga 300
tttggcctt ctcattataa taaagacctt cataacaagc ttcataggct catacaagga 360
anaaaaaagtg tagatggata ttataaagag atggagaatt ctttgagtag agccagtctt 420
aatgaagatc aat 433

<210> 203
<211> 240
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 203

acactcgnnng atatgnggac tntctctatt caacttnnnt gttcgctcta tcagtagaat 60
tcgttttagt ttccgtacac aatntcttt gtttcttcag tttacagtga gttagatatg 120
ttctaattat taatttcttt tcttttttt tcgaatattc catntaccat aaatagttgc 180
agaagcttct gtcacaccta taaaaacact ntatattga cttgacgaat ntatatgtat 240

<210> 204
<211> 437
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 204

gtgtccccca tcgaaatgan aaacctttat tcaaaccctt canagntagt gagaaggcta 60
aacgaaaaat aggaaactta gaaaagctaa atccttaact gaaggcgtag gtgacaatca 120

tagtgaatta ctaaacaaga atggtagttt acttaaggc attccagata cctcccaagc 180
ctcgaaaaat acttctaana tggtaacaag aagtaccttc aaatttaata atggtattaa 240
tgaagatagt gacccaaactc agatacacac ttggatagga cactatcaga aagatataat 300
ccaataattc aaaactgaaa cacctcaa atatatcag tcactgccct gcctctatag 360
agaagaggaa acatttaag ttagtgaata cattacgatg acataatgac aacgagatac 420
atatgatacc tcacata 437

<210> 205
<211> 505
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 205

ntgaccctga accctggatc tctaagtcac tctgaagatg cacgcntang aganaccata 60
aaaactaagg tagttctaa actaaaatca attgaggaag ctgcgccaag tatccccatt 120
gaaaaacctt tattcaaacc tttcaaagtt agtgagaagg ctaaacgaat aattaggaa 180
cttagaaaaa ctaaatcctt aattgaaggc gtaggtgata accatagtga attacttaac 240
aagattggta gtttacttaa agtcattccg gatactcccc aagccttgga aaatacttgc 300
aaaatggtaa caagaagtac ctacaaatta attaatgtta taattgaaga tagtgaccaa 360
agctcagata acacaactga gataggatca gtgtcagaaa tgaatataaa ttcaattaat 420
tccgagcact gggaaacacc ctccaaatta tattatcaac gtccaaactgg ccctgacctt 480
ctattataag aaagaggaga aaacg 505

<210> 206
<211> 424
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 206

tgctgactcg agagacaagg acaaccact ctaaagcatt tacgcatca gactcactgc 60
ttgaggggtt gtacatcatc cagaaagcga atcccaataa taccgactca agagacaagg 120
acaactcatg cttaaagcatt tatgccatta cgcttaatgc ttgaggggtt atacaccgta 180

caagatgaat attctattaa taatgactca ggagacgagg aagactcacc cttaagcatt 240
ttaggcacaa ggataaatgc ttgaagagtt gtacaccgct cgagatgagt attctggaga 300
tattgcctct agtgaggaga tgacttatcc ctanacattt atgcgacaag gctgaatgct 360
tgaggngtta tacgccattt atgatgaata tcccananat accgaccta gcaaaagaga 420
cacg 424

<210> 207
<211> 511
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 207

ntgaccctga aaccctcgna gcacctgaga taccctacag acgatgcgag ngatgcaagc 60
ttccncaaca tccaagtaac tctacattct aacaacacaa accatcacag ccaagaataac 120
agggcaaagg cagataactc tgcccaaacc accaacccaa atcacagctt ttctcactta 180
aagaccccag taacaattcc ttcgttccaa ttcgttaacc gttggatcga ctccaaaaga 240
ttactggaag tctctagtagtac ataagcctac attatgaccg ttggatcta cttagcaaaca 300
tccagaactc attctgaact actctgtcca cagccaatta cacacaagca ttgttctgca 360
cttgtgcaaa attctgctgc acaatttcac agcataaatac tgcacaaagt gcagatttcg 420
aataccacac ttccctctcat ccaatcttcg ccaaataaaa tactacaagt cccnatcatg 480
tatcaatcat gtctaaacca gagccaaagct g 511

<210> 208
<211> 257
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 208

actattccaa tacgctngga gcagcttaat agatctcaag ctgtgcaaattt attcagagct 60
aatagattga gctgagtcta aataatgaag gaacagttat tcaagttgc ttcttattcc 120
tccgggattc ggaaggacac atttgcatt cgatataatg attcacatcg tcatgttagtc 180
ttgaattatc gtacatctat attacgccccg ttctctattt ttgtacatga atatagacag 240

caacatgcaa tgaacag

257

<210> 209
<211> 488
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 209

accattgaa tcttgagacc atcgtnanc cagagatcct ctagagacga cgtcgacgca 60
tgcaagcttg ggctgggcct acctccatcc ctagagttag ccgtatgagg cgaaagctcc 120
acgtacggtt ntgaagccga gccttctag caatggngcc tagggaccga tatgtatgatt 180
ggtttaggta gggcgccgg cctactacgg gcacctgttag ggattagtgc ttgagaccgc 240
gatccacaaa agcatggac tcaccctta ctgagaatg aagagggaa tgacacgacg 300
tttcaagagc tatgcgaggg gtgaacaaa ctgcagaggg atcttcctga ccaggcgtga 360
tagagatgcc ctttattacc caactcatat tattcaaa tcctgcttg tgccactcag 420
tcttggcggg atccccctcc tttctttt ctctcaaccg gcgtccctt cctccacaaa 480
cgtgctcc 488

<210> 210
<211> 180
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 210

tgcattttc ttcaactcct gaagccatat catagcagac tcaagaacat nctgcattcc 60
anagatcact ttcatngaa tgcactatgt attggtagtc ttcccatcca ggaatggtag 120
atgcgttgaa ggtccgttcc cattcatctc taaaacactg aagttcagg gatcccacac 180

<210> 211
<211> 510
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 211

ggtgccctta ggcttgagac ccctgtagan cccgtgatac tttgcagacg acgcccangc 60
atgcaagcta ttagtgcata gtcacact tcgaaatgtat caaagtgggt ctgagaagag 120
gcaaatactga tcatacttgct ttgatacatg cgaacaaaca aagttggggc aaatacagag 180
ggtgatgatg aatgagaacc ccgagctgtg actgacattc ctatacagcc gagtttccca 240
ccaacccaac gatgtcatta ctcagtcgtat agccaacctt ctccttaccc accgcccagt 300
tatccacaga ggcacatccct ataatatcca cagagttgt cgttcgact ctcaatgacg 360
accatcatct ttagcacana cctagagcac caaccaagat atgaatttag cagcgagaaa 420
gcctgtagaa ttaatcccat tccagtgact tatgctgact tgctccata tctaattgtat 480
aactcaatgg tagccataac cccaaaccacg 510

<210> 212
<211> 425
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 212

acatttgctt gttgctcggt tgctgagttgt gtgatgagat cgtgttaggg ggttttaggg 60
aacttatatt tatatttagtt gagtgtaggt gctagtcctt gttttaggg gatattgttag 120
tctttgcagt taattctcggt cttgtagata ttaatcaata gcttacatataatgttagag 180
ataaacattt gcttataatggtag aaaaaggtag aagataatca aacctttagataatgttg 240
ggcttataaaa taattttttt aactgccaat agataagata ttcaaataca tttgaatatt 300
agtaggttag agataacctg tttgtttttttt agtccggctg gtnacgttca tccctcctct 360
cttttggcct gccttcattt tngttgcctc ctcctcatga ttccttctt ctacccctcc 420
ttccg 425

<210> 213
<211> 190
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 213

tgaagaccccg cacanacatt ngaaagaatn tcacattgtc tgctccacca tganacccccc 60

agatgtccaa gaggatcaca tatttctgaa ggctttcct cattcattag agggagtggc 120
aaaggactgg atgttattacc ttgctccaag gtccatcacg agctngatg accttaagag 180
agtattctta 190

<210> 214
<211> 224
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 214

atatanntt ttagtatta tatagaattt aatgttngaa tttaatttat cctgaggctn 60
tggaatgtga ttagcaatct aatgcataan agtattntat ttaggcgtg ggtgatcaag 120
tcaatctctg caaactgttc anattcaat gagattntgg gaaaagaatc tgagctggtg 180
gaatanatag ctgaggtcta atttcaatac aggaacatta tact 224

<210> 215
<211> 156
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 215

tcttggatgg ctactcggnng tataattttaga tagcagtaga tcctaaagac caagaaaaaa 60
atggctttac atgcccttgc ggtgtcttta cttacagaag ggatgccatt gngttatgt 120
atggccttgc caccttccaa agatgtatgc tagcta 156

<210> 216
<211> 195
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 216

aataccagaa gaatccaggt tctgaattcc atatctttagt tagaaggaaa ggttagcataa 60
tcaagtaaag caaaaaggaa ttgaatttga ctatcccatt tattgctcnn nnnnctngta 120
ttgtttgatt gtcttaact aattcttgc tttcaagcat ggcaattaca ttcacaactt 180
actcataatt tata 195

<210> 217
<211> 181
<212> DNA
<213> Glycine max

<400> 217

ttccaccatg gagatgcagc ggaagacaaa ggagaagagg taagaggcg cgccatccac 60
taggaaataa gccttggaaag aaggagcttc accaccaaga tgagccttgg ataagaagct 120
tggagatgtat gcttcaatgg aggaaaagaa agagggagag aaagagagag gggggagcac 180
g 181

<210> 218
<211> 274
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 218

gcagctgagc ganggtccaa tcattaccca ttatttacct tcntgnctta catttcttgc 60
accaacaact tcgtcacttt catanttggaa aggggtgcta catggaagtt gagggcagga 120
gaggaanaaa gatggatata attatgtgc gagtgactg aagggatngg tgtggattct 180
tccatcgtgg tcttcacata nttttggttt tgcatacaat acattgtgg attacgaaca 240
cctaaatcgg acgacacctgn tagctcttca cata 274

<210> 219
<211> 249
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 219

accatcacga ttatcgtctt cttgtcatt attggggta ccacctgagc cgccagatcc 60
ctacaccttt tgagcgtgtt ctttggaaaga tccgtcctcc tttgtgcaca tgctcatgag 120
gtgcacccata tccgaaacca tatcaagatt gtactgatac tgactaacac aggcaaccat 180
tatgtccttg caagaatgga ctcgtgaatg ctccaagtta gtgtaccang taacagctac 240
ctcagtaag 249

| | | |
|---|---------------------------|--|
| <210> | 220 | |
| <211> | 246 | |
| <212> | DNA | |
| <213> | Glycine max | |
| | | |
| <223> | unsure at all n locations | |
| <400> | 220 | |
| | | |
| gtatctngtt ntccatcgat gtgccatcat tntcttctat gttctanacc ctgtctcgca | 60 | |
| ccatttaatt attgatttgtt cttaattgtc aattaattag gcagttctat tatttgngcc | 120 | |
| cattcagcca atgtgatgct tttaatctaa tttcaggaat taatgaagaa ttgngcttga | 180 | |
| atctagcatt gngcttgaat ctagaattgn gctcgactt gaagagggca aactatatta | 240 | |
| ttctat | 246 | |
| | | |
| <210> | 221 | |
| <211> | 510 | |
| <212> | DNA | |
| <213> | Glycine max | |
| | | |
| <223> | unsure at all n locations | |
| <400> | 221 | |
| | | |
| gtactgcgcc tcgtacttca cancnctgta gtacccggca tccttagagt cgacctgcgg | 60 | |
| catgcaagct tgctctanat nttcattgtat gtttgtattt atgggaggag gtttatatgtc | 120 | |
| cattttgctt taagagtagc gtcccactgg taaaattaac tttccaaatg tttgccttcg | 180 | |
| caggaatggc cccgaggaag cttgcctcaa agaggtccag gaaggacaac ggcggcgaaa | 240 | |
| gaactatttc cgctccggag tacgacagtc accgctttag gagcgctgta caccagcagc | 300 | |
| gcttcgaagc catcaaggga tggtcgttgc tccgggagcg acgcgtccag ctcagggatg | 360 | |
| acgagtatac tgatttcag gagaaatag ggcgtcggcg atgggcacca ctggttactt | 420 | |
| ctatggncaa gtttgatcca gaaatagttc ctgagntta ttccaatgct tggcaacaga | 480 | |
| ggatggcgtg cgtgacatga gatcttggcg | 510 | |
| | | |
| <210> | 222 | |
| <211> | 207 | |
| <212> | DNA | |
| <213> | Glycine max | |
| | | |
| <223> | unsure at all n locations | |

<400> 222

tctcatgcac ttagtgtcca nataatatac ctatagttag aaatttatta tgttatggta 60
aaaatagggtt tggtaatct ctgtaaacca tanttgcata gntcttcaa ttaactgaaa 120
taatgagtct gtgagacatg acttaagttt aattctcaca gaatacactc ttngaaaatg 180
atgattagct ntaagtgtga ctaagtc 207

<210> 223

<211> 431

<212> DNA

<213> Glycine max

<400> 223

tctggtgat gagttatcga cagcgatgac tgcaacagt cctgagaatg tgttcaggac 60
tgattacgtg agatgcaatc tagaggattc tgcataatcga aagacttgg ttcatttcattg 120
cttggacta tgacctcgac tgcctgaga acattctccg aagacttcgg aaggatata 180
ttgaaatgag gtttcataac tccttcataac cattgataag gatctcgacg agttcttc 240
tctgtcttgc acgtcacgaa agatcctacg ctctggcgat cggtcaccag aaacaggc 300
tcggatccca ctggcgatca ttgctgtact attcatttc gtgcgtcaa gacgcttgg 360
tctgcattac cagaggtgtt atcagaacgc gtcattcgca tatctctgcc actgtcctct 420
tggtagagaa t 431

<210> 224

<211> 481

<212> DNA

<213> Glycine max

<400> 224

ttggcctgga gctggtcttgc acgaccccggttgcagcttgc cccagcatct ggatcacatt 60
ttacgacgca atttgggttgcatttc acgactaagt tctctctgc tccttactca 120
caccacgaac aatatagtttt agcgattaca ttgtaagagt attaggaact tgaaaactaa 180
gcacaggcta cactttaact aatggccttgc ttgctctgc cattcttggaa acctcagcct 240
actaattgaa acgttcgag catgttggta ttgactgccc atatggggc gttttctta 300
taaaggaaaa aataattggta tatataatcc gggggcttttgc atattgacag actgctccac 360

actaactttt ggtttatttg aaccctcctt tatagggctc aacacttgac acacgcccc 420
aataaggcct tgagaccaat atatgcttc ctctatccta tggtgccctta ttcctatac 480
g 481

<210> 225
<211> 277
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 225

acctagacta taaatagaag catgtgtaag actaggtgtg actgtgatga atgaaaagtct 60
tatgagatac acttcanagt tgcacttctt tgccctgtt attccttcaa ttgcgtgctc 120
cccccttctc tctntctttt cctccattaa agcatcctct tcaagcttct tatccaaggc 180
aattcttggg ggtgaagctc cttcttcctt ggcttattcc ctgtggatg gtgcctcccc 240
tatcctcttc tccttgcct tctatcgcaa cctaccc 277

<210> 226
<211> 279
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 226

acatcccact gagatgcact atgtaagtac cttnanaaa acagacttgt tgagaataag 60
ggagcttagca tctntagtaa gtgatccagn tgatntcaa gtcacccatg ggaagttgct 120
cagaattctt agagtagatg ttgaggaagg atgcctagag accctggttc agttctatga 180
cccgctctac cattgcttca catttccccga ttaccagctn gtcctcacac tngaaagatg 240
ctcctaccta gttggcttac ctgtgccaga caagatacc 279

<210> 227
<211> 220
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 227

tcgtcttatt canaaccnca accaattatg anatccncta tctcccactt cacacctcg 60

aacgcaccgt tcttatagag agaggcgctn tcacatcntt cttaggctgg gagagggaaat 120
gttcccatnt tttatgatac tccggngaac agatatccag tggagatgac ggngtggngc 180
ctgttagctca gaggattaga gcacgtggct acgaaccacg 220

<210> 228
<211> 379
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 228

gcagcttaga tgatagaggt ggagactcg gtcttctct tgcttgactt cttataatg 60
tggaggactt ggccctgcgac tcggctagat ggggctattt gcgtacacg cctttctcg 120
tacgccagct gttccgctac cgcatggta gccttcaata tttctagttt attctttga 180
cttaatcctt tacgattgca acaggaagga aattnaatt ttacaatagc atgatactgt 240
aatattnna gatatttata ttttagataa atactatttta gagtatacac caagtggatc 300
acagtgtgaa tgactgaatc agtgtacacg acaacagcaa gtgaaatcgg tgacaacagt 360
ctcacaaata tcggcacat 379

<210> 229
<211> 169
<212> DNA
<213> Glycine max

<400> 229

atacctgtaa tgctgcatacg ccacagaaaa tctcttcttc tgattaacac cacttcgtg 60
atcataagct tctggcattc tatactggca gttcaggaga acatatgagc cggtggaaag 120
aaacatatacg atttgtcaat ttacttataat atgcccataat ccctttgtat 169

<210> 230
<211> 497
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 230

nttaacccgt gagctttgag acccttcgag naccgtcatc cttggagaag agtctaggca 60

tgcgagctt tgtaattaga tacaaatgtt aatgtttatt tatataactg taaaaggcga 120
taagtgtggt gaaacaaaaa cctgaagtac catctcatta gtcagagtat atacaaggcat 180
gtacgggtga tagaaaaaaaaa aaagatgtt gaaggttgga tgacttctta gtgcataagag 240
tgttactcac ggtgatgatt tcttctcctg ccaccaaatg gcatatgtcg gcgatggaat 300
atatatgtta gaagggagcc tggcaatgat ggcattatat tattgcttc acatagatag 360
atatgtttta ataccttctg tcgacttccg tttatttatg aaatagtgg agggtgttat 420
ttcttgaatc cttgattttt tttaacgacc gaggtgacat tctatttttgc tctatagatt 480
ataaacacttt attaat 497

<210> 231
<211> 503
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 231

tggggtttc cggttggcc tatagtcctg gacntcaagt gctctgaggt atgcagatta 60
ccatcagacc acttncaggg tgctgtaact actttatatg gacttgatgg ggcctatgca 120
agttgaaagc cttggaggaa agaggtatgc ctatgccgt gaggatgatt tctgcagatt 180
tacctgggtc aactttatca gagagaaatc agacacctt gaagtattca aagagctgag 240
tctaagactt caaaagagaaa aagactgtgt catcaagaga attaagagtg accatggcag 300
agagatagaa aacggcatgt gtactgtatt atgtcatctg accgcattgc tcatgaggc 360
tctgcactca tcacaccaca actatatggc gtattgana tgctaacctc gactttgcga 420
gaagttgcct ggacttcttt ctccttaca acttcactca tttctgggtt ttcccttcca 480
ccgtctgctt tttgatgtat tcg 503

<210> 232
<211> 259
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 232

gtaatattat agccgatgct ctntctcgcc gtcatgcatt actttctatg ctgaaaacan 60

naatgattgg tcttgaatgt ttgaanaaca tgtatgaaaa tgatgaaact nttggagaaa 120
ttttttaaaa ttatgatatt ttttcagaan atggtttctt tagacatgaa ggcttcttt 180
tcaaagaaaa cannatgtgt gtgcctaaat tntctactag aaatttgctt gtttgtgaa 240
gcacatgaag gaggttaat 259

<210> 233
<211> 300
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 233

aatctcaata agtcaacaat gacattcgaa gaaatcgaaa tacgttatgg aggggatgta 60
ctatagccaa ttgttggaaag gcaatagaaa gaatngggaaa ccttaatcta atgctaagt 120
tagagagtga gagagagaat gggacttag taactcatga nagactntga agtctgaaca 180
agtttagtcaa ttgtttagca ctacatattg aagctttaaa tataaaanat atgtaaacat 240
ataaaacaaca atagtaattc taaagacatg tcacatcant gggcttatgg gttgggtcat 300

<210> 234
<211> 271
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 234

aagaatgtgt atgtttct tgatttcagg gttgtcatca tcaaananagg gaagattgta 60
gaagcaagct tcatgatgat gaatcaagtt gattcaagta gttctgtatga taacatagat 120
gatgacaaaaa agccaaaaga atgatntcaa gattgagtca acaaataaa gattaaattc 180
aagaatcaag agtcaagatt caagaataat caagatcaag aatcaagact catagattca 240
aanatcaaga gaagacttan tcaagataag t 271

<210> 235
<211> 212
<212> DNA
<213> Glycine max

<223> unsure at all n locations

<400> 235

gagagagaga ggngaggnga gcataatatt gaaggaggac aagagagaga gaagttgaac 60
tttgatatgt gtctcacaag actctcattc atcaaagtta caacaagtgt tacacatgct 120
tctatntata gccttagtag cttccttgag aagcttctt cataagctc ctgagaagt 180
tagagcttag ctacacacac ccctctaata ac 212

<210> 236

<211> 266

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 236

cacatcaaca tgctngaacc ttcactttt gtgagnaata agatttatnt ataagaatgg 60
ngaacaagga tcaaactcta gaccacacag ttatagaggc tccaattaca tgtcatgaac 120
ttgntactca aaagagctgg tgagttaaga catatggatg aatttatgtc tagcattcat 180
gttatggtgt tatatttgaa tattgaatat actatatgct ntgagaggaa nttaaacttaa 240
cttgctgttag aggaactgaa atatgg 266

<210> 237

<211> 338

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 237

acatctaact cagactaccg attcatgcc aataatatac gagacgcttg anattgaaca 60
acggaagctc tcgagagatt caaatggtca taactnttca catggatgtc caattcaagt 120
gcataatatt ctgagatgct ctaaattaa catggaagca caagggaaat tanaacggcc 180
ataacctata acaaggatgt ccgattcagg ccaataatat attgagacgc tcgatattga 240
acacttatgc tctcaagaga ttcanatngt catacatnt cactcgatgt tccgattcag 300
acgcataata taccaacatg ctcgatatta aacatcac 338

<210> 238

<211> 269

<212> DNA

<213> Glycine max

<223> unsure at all n locations
<400> 238

tgacccacgc gggtgtgaa gagacggcat gggcatctcc ctccttcctt cttgcccctg 60
atgccccat tcttnnngca tttagcggtta tggaagaaac gtaatcaaac ttccctctnt 120
tcaatccaac ctcgattttt tccccggcaa acaccagatc cgcaaagctg gacggcatgt 180
aacctactag cttctcatag tagaacactg gcagagtgtc taccatcatg gogatcatct 240
ctctctcaac catgggagga gctacttat 269

<210> 239

<211> 246

<212> DNA

<213> Glycine max

<223> unsure at all n locations
<400> 239

ctacactgac tcccgagttt tctaagggtt agtggcanac atatatcaag ccaaggaggc 60
agtattgctc aagtactatc atgctgcana aacccttatt gtatgtactnta atcgcttcaa 120
gatgtaccat atacgggggg agaacatcac cagagcagac ttgctctcca agttggctag 180
cactaagaga gctggacatc ttaagaccat tatgcaagag acactccaag cacctaccat 240
agacac 246

<210> 240

<211> 271

<212> DNA

<213> Glycine max

<223> unsure at all n locations
<400> 240

acaacacctaa gatttatcat atacttatca agagaatcaa tccttaagag agactntcca 60
nattcattgg aagcactgag aatcttgaca aactgttaat atacagttaga tgtccttcta 120
acagatctgg acatggatata gaaggagata cttatgttca tgataaggaa actaccaaata 180
gttatttctg tgaaaaggat caatgttngg tcaaagatcg cattggagac ccttctacca 240
ttgtcatgac ttgagtgccct tcaatcacta t 271

<210> 241
<211> 425
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 241

gatcgccgt attcctggc cgacgcgac tgtcattttt ttcgatcaat atcggtgaat 60
aataactttt ttgccgaggt gggctaattt tttcctggcc gaataaatgg gaacatgcca 120
gtttcggcg aaacgaaaca tcgggtgagc tcgcacgaaa aaaccttagcc cacctacatt 180
gtaagtttt tatgcaacac cgaaacaaga aaacttcccc tgccgtaaga aaaaacatta 240
tcggccagcg agcattttt tttaaaaaaa attgcgcaat gtcggctgaa aaatatcagt 300
cgngccatt tcacgaccga tgtcggctat tttggttct attcaatccc tgaatgaaat 360
ttgcatgatg tcgattaaga aatgttngat ccgcgtgatc cggtgatgct tttttttaga 420
actcn 425

<210> 242
<211> 253
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 242

tcttctcaat gtccgttaag tcttatggac ttcttgccata taaaggataa gcctttctt 60
cctgcctctc attntttgct tcctgcacgg ntgcacctcc aatatnntta atatccnctt 120
attacccttc tctgcttctt ctgttggct tcttctccct tctcaacact atggcagaga 180
tcctatgcaa ctgaaccana ccanaatctt ccatcaagaa caaaccanat tngaaaaacc 240
acaatggaca aac 253

<210> 243
<211> 393
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 243

tgcttcttctt agaagcaatc gccttctgga ggaattttctt ggaaggccca agtgggccta 60

gttgcttattt gcaccccat ttttactaaa tacaccactt gctcttttc ggagattcc 120
ctccgcacca aaccccaact ctcccctgt ctcgtcctct atcacgtcag ccctcaaacc 180
tttcttggtg tcctcctcct ccctcatccc aattactacc cctacccaac tacaaacatt 240
tacctcacgt cgcgttact tcacatacct ccccgccct gcgcattcct ctcctaccct 300
tcgcccaccta gtccccattt tccctctntc tccctctcac ttccctgcttc cacttacccc 360
ccccctcttc cttctctcct ctctcctctc ccc 393

<210> 244
<211> 416
<212> DNA
<213> Glycine max

<400> 244

tattttcctg aatcgacat ccgagtgaaa agttatgacc attggaattt ctcgagagct 60
tccgctgttc aaatttgaga gtctcgatat attatgtccc caaatcgac atccgagaga 120
aaagttatga ccaattgaat ttctcgagag cctacgtcgt tcaatatcga gcgtctccaa 180
tattatgcac ctgaatcgga catccgagtg acaagttata accaatcgta tttctcgcg 240
gctttaggcg cgcatatac tacacctcca tacatatatt ccactgactc gtcatcctac 300
actcaaacct attatcatct ccatatctt ttgataaagg ttccatccc atactcactc 360
ctatctctcc ttcatctcg ccatcagttc ttatttctt tccactcgat ttatcc 416

<210> 245
<211> 222
<212> DNA
<213> Glycine max

<400> 245

catgcttcta tcttgagatt atgacacatg ctaattcgga ggagatgatg attcgcgaga 60
tgtatcgagt catctcgat attgtggttg gtactctctg aacaggtcat taagcgaact 120
tagcatagtt agctctctt tgcttagagga caagcaaaac tatacatact ggggagttt 180
atcattgacg tacataagtg gattatgcaa ttaagacata ta 222

<210> 246
<211> 429
<212> DNA

<213> Glycine max

<223> unsure at all n locations
<400> 246

gggatgtgag ctctttaga cttttcttg caggaatgta agcttatcc aagaganatg 60
tatggtcatt gggtgtcatg tttcaatta gctccatagc ttcatttaggt gtttttaact 120
ttatcttacc tccaacggaa gcgtctagaa gttgcttcga gtgaggcgt aagccatcaa 180
taaaaatatt tagttgtatt ggctcaactaa atccatgcat tggagtctga cagagtaaac 240
cgtggaagca atctagagct tcgctgagtg attcgtagtggactgatga aatgaaaaga 300
tatctacctt tccttgcgt tgcttgatt ctggaaagta tttcttcaag aatttatcca 360
caacctcttc ccaagttctc aagttatttc cttaaacgaa gtgaaaccat ctcttatctc 420
actggcagg 429

<210> 247
<211> 215
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 247

tctcaactacc ctgttctgc cataacttaa gcctatagat aattcttac cccatgtngt 60
ggtcctaaag aatagtataa tcatcacata ctaggtatcg tttggaatct tttaaatgag 120
aaagaggata ctatctcgat aatcaatata taatcataat tcatgagttt caatctaattc 180
aatataatca taattcatgc taatcaatat atact 215

<210> 248
<211> 510
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 248

atgacactga gactcttgn actnccctag aatcctctag agtcgatgct gaggcatgca 60
agcggtggca aatggagagg aatattttgt ttctatatgg agacgacgaa gagagcaagg 120
tgttggaggt ttctgacaaa attgaaagga tttgatttga tcgcaagcaa taggtgtgg 180
agtaagggtta ggggacgaaag gattggccgt gtgataggag ggcaacgaaac aaatttccaa 240

ggcaacgaaa aatgatttac attttttatt ntgttcgcta aatgttcat aagtcttact 300
aaagtgaatg caagcaaaag aaacaatata ataaaaagga taattaccat ttttagtcatc 360
cttcaaattt caagggttttgc tcccatctt ccatgacaag aacaatattc atattaattc 420
acgtgtctct caatacgtcc ctacccttaa aactcttca ttccttcacc catcctgtca 480
ctctctctct tttccatcta ttatgtgtcg 510

<210> 249
<211> 239
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 249

atggtgtctt ggagcgtgct tgtgattntn tctttcgcca tgctgcacaa ctntcaggtg 60
ttcccttgag aatggtgag agaagcagaa ggcagttccc tcttagaana gcccggtatg 120
ctgcagaaga catgctctct gggctactca nagccaaggg tggatggattc atgacattga 180
tcgaaaatgg taaatggatg tgtgacgagg cacctcacag tggaaatgaa tatgtaaac 239

<210> 250
<211> 255
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 250

cgtggngca caacaagctn tccacatcca caatgcgcgc ataaacccac catccnctgg 60
tgccccaccc caactgagct cacgtactcc cacgttagccc atctcctcgt ttctctcaac 120
accgggtccc catcaatcct ctcaagcttc cacaacatcc aagcanaaca acgttcaaac 180
agcacaagct atcacagcca agcaaaacag agcagaggca gataactctg ctcaacacat 240
caaccaaaat cacag 255

<210> 251
<211> 230
<212> DNA
<213> Glycine max

<223> unsure at all n locations

<400> 251

aatttctggg tccaaaataa tgtccaatan aaatgcgact catcaactgtc aacgtaaaca 60
attaacaatc aacatcta at ttcggggcta gagactaaaa tagcgagatt acaanaaaat 120
ggaagactca attgataaat taaattacag gggAACCAA ttgtgagcta naatggaaga 180
ctcaatccga cggcctcagc taccgtttcg aactcattcc ggacaccaat 230

<210> 252

<211> 224

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 252

catctacaac agacccctt aacctcagca gtatattcat ccacaacaga ataattatga 60
cctctccagc aacaggtaca atcccgagtg gagaatcatc ccaaccttag atggcgaat 120
ccttcacaac aacagcagca acaacaacaa ccttaatttc anaatgctgc tggcccaagc 180
agaccatacg ttcccccacc aatccagcaa caacaacagc aaca 224

<210> 253

<211> 426

<212> DNA

<213> Glycine max

<400> 253

caggctccac cagttctagt gaacgtgctg cccataccat tggcgcttcc aatgcggcgc 60
atagatgtga tcgcggcaat tgaacccaaa gcttccatg ggcatcgcta catcctagac 120
gctatcgatt acttcactaa gagggagaa gccggttcat atgctaccgt gactagaaat 180
gtggcggta ggatcataat aaaggagata atttgaaat atgggctgcc gagctatatc 240
atcactaaca acgccgccaa cttgaataat aagatgacga atgagttgtg tgggtattcc 300
aagagaccac accataattt gactccttat cgacccaaga tgaatgatgg agttgacgcc 360
gctcacaaga atactaagat gatcatctta gagatgacag cgacatacaa aggattgcac 420
gagacg 426

<210> 254

<211> 505

<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 254

ttgaccctg agnnttgaga ccccgtagn atcgncgatn ctggagaa gaantctagg 60
catgcaagct ttatggaa gtatagagca tganagtgtt ctgataccat taactagtca 120
atagggtctt gagcagggtg aggacatcca tactatattt ggaaagaccc aaaagaagga 180
aaaaaaagagt aaaacttgca tatggaagaa gaggtcgata ttgttgatc ttccatattg 240
gtttgatcta gatgtcagac attgttagcaa tggatgcac gcggagaaaa atgtgtgtgg 300
tagtgtcatt ggcacacttc ttaacattca aggcattgaca aatgaggcgt agaacactcg 360
acacgatctt attaatatgt ggatctgaga ccacttactt cctagggttg acggtaaat 420
gttatccctc cttccacta gtgatacttt gttcaataac acattaactt tactccatc 480
ttattctacg ctcccttctc ctccc 505

<210> 255
<211> 432
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 255

gttgattact attttttaac atcgattta gcgtcaaccg atataaaaag tgcttacaa 60
caccgatttt cattagaacc gatgtaaaaa gtgcttaca acatcaattt tcactagaac 120
cgatattaat gtattnntg cataattaaa aaaatatttt gttcacaag aaccatttc 180
ttcgtgatcc attttttaa aaacataatc ctgtgaccat gaagaacaaa actaagacac 240
tataatatta atataataaa tgcaacatga agcagctagc gaacttcaaa aaaattattg 300
cgaaccaatg atttgatgtt ccagtagcaa ctggaccct tggatgctc gtattgctta 360
tagcatccta naagaataac aattattaatgt tgcattgggt ggttccaat tatccctgaa 420
aatatgtcag ct 432

<210> 256
<211> 206
<212> DNA
<213> Glycine max

| | | | | | | |
|------------|---------------------------|-------------|------------|-------------|-------------|-----|
| <223> | unsure at all n locations | | | | | |
| <400> | 256 | | | | | |
| gcataatcca | taatgctgca | naacannata | cacgcataaa | taaaactctta | atcgacatat | 60 |
| ggagtacata | ggaanatcct | ttcttaatag | ttccagcctc | catagacaat | aacgttgctt | 120 |
| atgaattaag | aacaacaata | aaactagtct | aatatgtga | aacacataat | ctgaagaaag | 180 |
| aatagctata | ttcatcaaca | gcaata | | | | 206 |
| <210> | 257 | | | | | |
| <211> | 221 | | | | | |
| <212> | DNA | | | | | |
| <213> | Glycine max | | | | | |
| <223> | unsure at all n locations | | | | | |
| <400> | 257 | | | | | |
| tatagtaaca | ggcanacgga | aggattccac | acatactaag | ctttcattgg | tgatgacgaa | 60 |
| tttgttgaat | ntgatgagaa | cctcattgca | atcttcaagc | acatattcaa | cctttatgga | 120 |
| gaatattctg | atagtgccaa | aaagaagata | gccaagaact | ctattgcana | atcttgcctg | 180 |
| cagaaggaa | aaatganatg | aataacataa | cataacatta | a | | 221 |
| <210> | 258 | | | | | |
| <211> | 478 | | | | | |
| <212> | DNA | | | | | |
| <213> | Glycine max | | | | | |
| <223> | unsure at all n locations | | | | | |
| <400> | 258 | | | | | |
| gcctcgccaa | gcattccat | tgaagaacca | ttattcanac | ctttcanagt | tagtgagaag | 60 |
| gctaaaagaa | aaatttaggaa | acttagaana | actaaatcct | taattgaagg | agtaggcgt | 120 |
| aatcatggtg | aattactaaa | caaatnggtaa | gttacttaa | ggcatcccc | gaaactccnc | 180 |
| aaacttcaga | taatacttcc | aaaatggtaa | caagaagtac | ttncanataa | ttaatgtatt | 240 |
| aatgaagaat | gtgaccaacc | tcaaaaatca | agtggataga | tcagttcaga | aagaatataaa | 300 |
| tcattaatcc | aacactggaa | acacctctaa | tatataaac | gccaactgcc | tgactttata | 360 |
| gagaagaggg | aaacatntag | agtttagtcaa | cacattatga | tggactgatg | caacgagata | 420 |
| catatgatcc | tcacacatga | ctgagtcact | acaactcatg | atgtaaaaca | tatataatt | 478 |

<210> 259
<211> 507
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 259

aggatgact cttttagacc atcgatccctc agagacgatc tcgcaggcat gccagctaa 60
tgagtatgga agcgtcacaa ataacttctt agtttttat aatctaata ggaaccagaa 120
tctcgagggt gtgaagacac ccactacctg cgatgagttc ccgggtggag tactctggct 180
attgtaaatg gtatagtcat gatagcttcc attgttttagc actatcacgt ttacattat 240
ggagaaggcg cagacagagc ttatTTTATT gctgacgtgc acgcccaggc gcagcactct 300
cgagagatc ttttaacagt atacgcttct gcgccatctg tgctatgata attcgaggtg 360
aaccagaata ttcttatcat gctgctcaag aacaatgcta gagtttatat tctgagctat 420
ctaananacat gggacattcg ctactccgtg aaataacaac cggccagaat tctgcacgat 480
accttctcat ttattataat cccaaag 507

<210> 260
<211> 290
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 260

gtttaagtct tctaaactgc ctatctatat tccatagcct ataacaactt ccgtgtgccc 60
atcggtttgt gggtgacaag ccggtgacaa taacaattta atgccaact tgctccacaa 120
agtccctccan anatggctta agaacttaga gtccctatca ctaacaatgc tccttggcat 180
accatggagt ctcacaatct cttgaataa cacatcagcc acatggaaag catcatcaac 240
tttcttacat ggaatataat gagccattnt agaacaccta tcaacaacca 290

<210> 261
<211> 428
<212> DNA
<213> Glycine max

<400> 261

gattggtgaa tcttcctgct tttattggtg accacagagt ggtacctgga gatatgtcgc 60
ggcggtcacg agaccttggg gacgtcaggt ggggtgctat tgcccaaaac caagcttgcac 120
caatcccgac ccaacccggg catagtcggt cagtgagaac ctgtgatgta cctaaacagg 180
cgagctcctg gcagtcaca gataaaagga acaaagacca caaaggcaagg aggcttgtgg 240
tagctggcca gctgtgaaac ttgattgata tgtgagatat ggtctctggt aatcgattac 300
caagggtgga gaatcgatta caaggcttat aaatgaagac aggttgctaa catggctct 360
ggtaatccat taccatatgt tgtacccgcc aacaggctcg gacactgatt cgagaactat 420
aggacccg 428

<210> 262
<211> 421
<212> DNA
<213> Glycine max

<400> 262

taacacttta acgtgcatac gtttagcaggg tggtaattaa tacacactgc ttccgtactg 60
ctaaaccaga acaaaagatg attaatggta gatgtttttt attatcgata aatattaatt 120
tattaatttt tggtaaaaat atagctcaca agagtaatat ccaattcata tatggttcta 180
tattgtgatc ccaaaagatg cacgacattt tttttcttc tttccacatg aactcatata 240
tcatcctaattt ctacaatggt ataagtgaag attgtaccat acctcatata atactactga 300
gatatgatca cgttcacaaa tgaatttcat tcccatttat tctctactct ggataacatc 360
ctccaaactat attatcaacg cccacctgac cctgaccttt ttttagaaaaa gaggtgtaac 420
g 421

<210> 263
<211> 493
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 263

ttgccctgac cctggacctc gagaacgtga tacttacact cacctgaaga tgcagcttga 60
gaatatgtct atactggatt aaccnattat agccttatca taaacgacta cacaattgct 120
ctttagacaa cgactgattt attcaagaga ctgtacttta atcgattacc atgccccatata 180

atcgattact tcccttgta tgcgtgtc agaagcgaac aagaacactt taattgatta 240
ctttgagttt ctaagtgatt acatagtcct tatgttcttt ccaatttcg agaagaacgg 300
tttaatcgat taccaagata atctaattcgat tacatcattt aattgagcga ttaccttgc 360
gactcaactg ataacagacc ggtgttaggtg ttttctctat aaacaaccaa ctgtgctat 420
gtataacaac acaacaattt gatctctagc agagcctgca tcacttgc 480
aagaaaagaag cat 493

<210> 264
<211> 509
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 264

ttgcccttgt anccttgnntn acntccggag nnccgctgat ggtatgcagt cgatctcggg 60
gcatgcnaga ctaagcgcat agttggggca tttatcacta ttacctgtat cacataccctg 120
atccgattat gacttgtact gtaaagaatc acgagtgatt gatattatga tcctcgccac 180
cttggttttt tacgaacaca tcataacact tatagagntt gtactacaca cagcacactc 240
tctgatttag ggataggtgc atgcttgaac tatgattgct gggattgctc gattgttagtt 300
tgagctcttt tacgctgtaa actcgccgtg acttgagccc aacatcactt acatacactc 360
ttctatatgc tcttactcgc ttcaaattgtat atatgattt gaggcgactt taactcgct 420
gtgacttgcc catgacaact catatgactc ttctcaaattc aatctacggc cctatgttat 480
taccaatgtg agttgatatc atcctaagg 509

<210> 265
<211> 485
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 265

atganccttg agccccctcg agngccggg atccttagag tcgaccggcg gcatgcaagg 60
tnngactcat gagcggcccg tttatgactt attgaattga gccttgcgt gttacttagtg 120
ctattactta tggcaccgat gagagcgcac cgactactca ctcaacagcc tattgtaaat 180

ggacactgac taaacccatt cctaacacac cgttctatag cgcgacgagt ggcgtgagag 240
acttacttac tccagcctac tcttcacaaca ccaagactta cagagaattt catttcctat 300
ctccctcatt aagtgtatta cactagtac ctcttatgcg atggttgac tagaaaaaga 360
gcttaactcg tgccacactg acgactatga gcagttacatt agagcgatgc tgttaataac 420
agtccaccc tattttcctt gtcaagacaa atctacatgg acctctgtta gggggtttcc 480
caacc 485

<210> 266
<211> 435
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 266

ggaaaactta ggcaatagac actagtatct tagcaaaatt attattatta ttattatttt 60
tactttaaag aatagctagt cattgaaatt ttctcccccc ccccctttt ttcttaaaat 120
tttatattat tatctatata gttttcccta cacacacaca cacacatata tatgtatagg 180
tatataattg tacatttatt tctggcaaatt caagaagcta atttgcac aaaatcactc 240
tagaatagcc ttaacatgag tccaaaacaa catttaagca caaatttaaa tcctcttact 300
tgcttcaaatt ttaataagaa atgggttgta cttaaactcc acaagttact tgcccatcaa 360
caactcanat gactnttctc aattcacaca acaaccctat ggtaaatacc aatttgagtt 420
gatatcatcc taagg 435

<210> 267
<211> 200
<212> DNA
<213> Glycine max

<400> 267

atgaactcta tggacttagta ctcattgtgc tggatacata cttgataatt gttattgtta 60
gcagcacctt ctctgttcat ttagtttcat atattcatcc ttagacatata atgatcttg 120
taggctgaca tataatatgtt acatggaggg aatgaaagcc agcgcaattt cactcactgc 180
tcttagcagaa tatacaggtg 200

<210> 268
<211> 272
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 268

gctcttaact gcacaaggct cttaatatgt gaagagtatc cttgtgtaac ctgcacncg 60
cgaagacact gacanagact tatcttcttc ttattggaca cagtatggca ggctggcggc 120
aagtaaatat tcataccatc agaccttggaa tgcaactgtg atcgatgcc catatcagct 180
agatcttaac gggatttcaa gccatccttc gtctgcctt gaatggtaag gagcgtacca 240
atcacactgt cacaaacatt gttctccaca tg 272

<210> 269
<211> 451
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 269

cacgcctgtt catntgactt tgccaccaca agtgggtgac tctctatcgt actcacgctc 60
caggacgcgt ctccctgcaca gaatgcagaa ctctctcaact gcgcaaaagg agtgtataga 120
cttttcttca tatatcgagg cccgcattgc tcttcgtgac actattgacc ataaaggcatt 180
cttgcacatg catgatatga aattctgatc ctaatcatat gcctcctcan gttcatctcc 240
aagtacagtt gacagtgtgt gtttcttatg cacccttaac tattgcaaattt atgatgccgg 300
ccactataca tatcacgctc tctacaccag aggacctttt ctatcttgaa cacttcataat 360
attgcgccca tgacctatcg catccccact attacgcctt gaggaccaca atgcttacca 420
atactcctct ttctcattac tacacaaacc g 451

<210> 270
<211> 262
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 270

gaactcttaa gagaaaaatgc attagaatct ctgttgcgtatga atcttagann atattggacc 60

```
gtgataagtt cgccanaaaga gtcagaanag acacanaaca agtaaagtat ttatagatnt 120  
agcaganaga aatagatcca aacgattagt aatcactggt atttgattaa ttngatcata 180  
ataccttgt tntgcattnt caaaatcatg gtaatcgatt acaatatgtg gtaatcgatt 240  
atctcannat aacatagatc at 262
```

| | |
|-------|-------------|
| <210> | 271 |
| <211> | 146 |
| <212> | DNA |
| <213> | Glycine max |

<223> unsure at all n locations
<400> 271

cttcattttc tccatgtatt tcctcacatg tcttatgtga atggtgtaac atgattctnt 60
agaatttcca ccaatttaac ttgctataga agcttagatn gatttcctct agttcanatt 120
tcttggtctt gntttgaac catgaa 146

| | |
|-------|-------------|
| <210> | 272 |
| <211> | 99 |
| <212> | DNA |
| <213> | Glycine max |

<400> 272

cgtatccgg tgagagtgtg atccttata tttgagagaa acagctatca tttagcattg 60
atttttgcat gaatctctga agtatggact gaatgcattg 99

| | |
|-------|-------------|
| <210> | 273 |
| <211> | 227 |
| <212> | DNA |
| <213> | Glycine max |

<400> 273

ctctctgacc tggaaatttc cgttcaactt attgaccatt agttcgagt cgatgaacac 60
gtatcttgta ttatttatat tatttgtgag cggaaaggct attgttaatg ctgttcaag 120
taccacccca tatcaggatt cctataagat tctcgctcca catagtgtt tactggatgg 180
gccattgaca aatgttaaacac cctctaccct cacacataac qaataaa 227

<210> 274

| | |
|--|---------------------------|
| <211> | 418 |
| <212> | DNA |
| <213> | Glycine max |
| | |
| <400> | 274 |
| | |
| accaattaaa gagttcactt ccaaacatcc cagccttcg tattggtaca attgaaacat | 60 |
| tgagaaaataa tcaagacttg tgtggaaatg tctctggctt ggaaccatgc cccaaaagcaa | 120 |
| gtaaaaaaaaatc tcaaaaatcat aagactaaca aagtcatatt ggtattttta cccggttggtt | 180 |
| tgggtacttt aatattggca ttatggcct ttggagtctc atatcgctt tgtcgaagct | 240 |
| cagagacaaa agaacaccag gatgaaaaac caccaggcca aaatctattt gtgatatgg | 300 |
| gttttgcgtgg aaaaatggtg tatgagaaca tagttgacac cacataagag ttcgacaata | 360 |
| aacatctcat tggagttgga ggacaatgaa gtgtttacaa agaagaaatt gacatact | 418 |
| | |
| <210> | 275 |
| <211> | 508 |
| <212> | DNA |
| <213> | Glycine max |
| | |
| <223> | unsure at all n locations |
| <400> | 275 |
| | |
| atgaaccatg aancttgcata ccctcgagan cggcgaact ctgagacgac ccgcggcatg | 60 |
| caagcttgtt gagatacctg gcgaatctca tttttctat ataaaggtaa tagaactaaa | 120 |
| tcatataaat aaaaggcaga caaaaatgtcc attgtaaagc atgaccacat caatttcatt | 180 |
| tgcattagtc ttgaagacat aaagattcta catgttgcaa gaataagaat gtaacagtag | 240 |
| aacatcccat ttctcagaga atgtttacct attccaaaat atgggtgtct tcaatgctaa | 300 |
| gcaatgtaga ccatgctcat tgtgccatc catttgaatt tcactcaagt catcaccaac | 360 |
| gttaacaagt tctctagagc acctgactat cagacgacaa agatctttg ataaagaaac | 420 |
| ctcgtgacaa gtaacaacat cagctccac aaggtgaagg ttaatacttt tcctcaatga | 480 |
| ctcattcatg catgatgcat acacacccg | 508 |
| | |
| <210> | 276 |
| <211> | 437 |
| <212> | DNA |
| <213> | Glycine max |
| | |
| <400> | 276 |

cttcaaacac ttgtgtaatc gattacgatc aacctgtaat caattaaaac aaagagttt 60
aactatagag gaaatcttct aacttagaa ctttcttct aaccctaca tggatgatgc 120
tgatgcacat atgatatgat agagactaag atgcaacaca caatataaca atcaatacaa 180
atgccactca agagagttgg gcatgtaaaa aataaaacat cttaaagctc ttcttcaagg 240
ttcaaggcta acgtttcatg ttgctccctcc tatctctaac aatatttca tggcacaaaa 300
catatatata tatatatata tatatatata tatatatata tatatatata tatatatata 360
tatatatata tatatatata tatatatata aaagtgaatg atatgtttt cacatagaag 420
gcgttccacc acaatag 437

<210> 277
<211> 519
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 277

cgttgtgann ccatcgatna actgacactc tataatactc aagctggca gcanacgtat 60
gtatgattnt aagttggctt gggagttgggtt gctgacattc ctttcttagtt gcacctatta 120
ctaataattta tgtttgaatt cgcatgattc tattttgatt ttaacatgac ttatgtggga 180
gcagggatga tttcccccca ccttggatat tcgtcttgc tccttggagc cattctctcc 240
ttcggagtga tatggccgct cattgatcgt cgcaagggag attggttccc taccaattta 300
gatgagagca tcatgagagc tttgtacggc gtcgaggctt ttctaacagc tgctctcattc 360
ctcgggtatg gcttatacaa ctttgtcaag atttttagttt tctcaatcct tagcgtacat 420
gaaataacta agaaccgtgg aaatggatag acgtacatgt caattccac tctcttgtct 480
gatatttagtc atgaaaattt cgtcacaatg ctcattccg 519

<210> 278
<211> 453
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 278

tatctccgac agccaatggg tgagtctcgt ccaggtagtc ccgaaaaaga ctggcctcac 60

agtgatcaga aatgagaagg aggagctgat tcctactcggtgcagaaca gttggagagt 120
ctgcattgac tataggaggc tgaaccaggt taccaaaaag gaccatttc ccctgccatt 180
cattgaccag atgcttgaac gcctggcagg taaatcccac tactgttcc ttgatggtn 240
ttctggttat atgcaaatta ctattgctcc tgaggatcag gaaaagacca cattcacctg 300
cccccttcggc acttttgctt ataggaggat gcctttcggc ctgtgcaatg cccctggta 360
cttccagcga tgcataata gtatttcag tggatttta aaaaattgca tagaggtgtt 420
tatggatgat ttcactgtat atggatcctc tttn 453

<210> 279
<211> 265
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 279

taacgatttc taattatgtg ggccattaaag tctatcatat gctaacaata gccgagaagc 60
ccatgaatct ctcggnggc ggagtaggtg tctgccattg ccttggcattt ggctaaacaag 120
cggnngaagtt cttgactccc gttcaaggta agagcaaacc gatccatcaa catggttgcc 180
tcttagtgta aagagtcgat cacccttcct ctagcctctn tttccgtgta tacttgagca 240
tactcatccg cgattctatg ctctgt 265

<210> 280
<211> 456
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 280

naggtttt gatgataaca atgatgacaa caaaatatga tgaaaaaaaaa gctcaagtga 60
atcaaagaac atctcaagag aatcaagaac aagtcaagag ttcaagaatc aagaagaattt 120
caagattcaa gaagaaagcc tacaacaag aatcaagattt caagattcaa gatctcaaga 180
atcaagatca agattcaaga ctcaagattc aagaatgaag aaaagactca atcaagataa 240
gtattaaaaa gtttttcaa aactttgaat agcacatgag ttttgacaa aacctttacc 300
aaagagtttt tactctctgg taatcgatta ccatattgtt gtaatcaattt accagtagca 360

naatgagttt gaaaatgtnt tcaaactgaa tntacaacat tccaaatatt tc当地aaaggc 420
tggaatcgat tacaatgttg tggtatatcgat ttaccg 456

<210> 281
<211> 264
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 281

gtatcacaat atatgttagct acttcccttg agaaaaaatt aatggaaaat taattttagt 60
ntatatnnta cttcaatttt ttggcaatgg ccggattgaa aaatatttaa ttgaataagg 120
gtgttatatt gttgaatntg tcaataatac atgtgttgaa gttattttgg tttttttac 180
ttgattactt agtacccttc tctatattaa cgtttattat anaataagtn gtttataact 240
atgactaattt gttcatattt aaat 264

<210> 282
<211> 246
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 282

agcaattggc aatttatact ggaatattct aaattacaaa tgtntataag aaacaaacct 60
gcaaattgact agctacttgc tctcttatca aagcaggat attcatagct tcaacaatcc 120
tctttggctt tgcttctgaa gaataatcaa gggagaagat taattacaat tatcatagaa 180
ttaagccata tacatgttnt taaataccaa ctaagaggaa tacanatggt gaagaataat 240
catact 246

<210> 283
<211> 457
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 283

taaaaactat aaaatgcaag gttaataaaa tgacattcag tttgtagata tgggggtct 60

ttctaaacaaa caagctgatg catagaaaata tatttctcta atcaatcggt ctcttggtt 120
ctatgttgta gcctaaatta ctaaacccctc gatccctcggt caggatgaat atccaagctt 180
tgtccgcaga tcacctcattt aagactacac ctgatttaga cagccctt aggtatagac 240
taacttaaac ttagtntcat ccgcagatcc cttatgttaag actagactca cttagtgc 300
ttaccaaagt taagcctatt taagccaaag ctttgaccgc atatccttat gttagactagg 360
ccaacctaac cagctttatg tacagcatat ttaaaccaac cttacctcgac aatccctcat 420
gaaggcttaag ttaatcctgt tcatcaattc taggcag 457

<210> 284
<211> 429
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 284

gctcaactaca agccttaggt gataaaccat gatattacca tatttccttaan ggaatttgg 60
gctttggaaat tgtnntggaa ataagtgtgt ggggtctatg gttcatagga acacatgcct 120
tgtnactat gttcatgtat gtattntggg ccatacttga tgtacattgc atattggcta 180
aatgttggac atgctgaatg aaatgttggtt tctcataggt aaaagaaaac aataaagaga 240
acagcaatac agttgagtga ataagatctt aaatggcaca agaatgtga gactcttggt 300
tctactcttc atgtctaatt ctatcttgcac tcttttattt cgttagtgttt taatatgcac 360
tatcccttng ctctcttattt tttggattac cactattcat attctcatac ctgccttgc 420
ccatacacc 429

<210> 285
<211> 544
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 285

cgactgattt atttgattca gtactcgatc acgctgatcc tttatcactc acctgcgcgc 60
atgcaagctg gcttctacaa ccctgtcaaa cttaagtgtg ctcgnagnng gactgacaac 120
ccacgttagag tctgaaaacg agagtcttca cagtgacacag tcatgactgt gatgagccta 180

gtcctactcg ggggcagatc atatggagag tctgcttga ctattaggat gctgcaccac 240
ggttaccata aaaggaccat tttctcctgc cattcattga ccagatgctt gaccgcctgg 300
cacgtcaatc tcactactgt ctccttgatg gtttctctgg ttatatgcag attactattg 360
ctcctgagga tcagggatag accacattca cctgcccctt cgccactttt gcttataaga 420
ggatgccctt tcgcctgtgc catgcccctg gtcctctcag ccatgcatga ttactatTTT 480
cagtgatttt atagaacatt gcataatggatg gttatggatg atctctctgt tttcgatcct 540
cttt 544

<210> 286
<211> 451
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 286

acanaatcta tgtatccaaa acccctcaat ttaatggatt ntcaagttt gagaagtgaa 60
attggaaatg ggataaattt ggagcaaact ctcacccac acaagtctat aacatcaatt 120
taaaacttgtt caaactggat ttacacctaa aatttcactg aatcaaaatt tgactcctca 180
acccccaatt ttaccctaga aatggcttggat ttgcattttt ggtcattttt ttttctctt 240
agcacagccc anactttctc ataagtccta aatgacattt caagcttagga ttaactcact 300
ttaacctcca aataccacta aatccagatt tggccttcca actctaaaaa attcactctt 360
tntccactca taacaccata atctcacctt ctaacccttg gtaattctac ccttatctct 420
aacagtttcc ataacaattt caccacaaca t 451

<210> 287
<211> 54
<212> DNA
<213> Glycine max
<400> 287

gtgaaagggc tagtgatgtg cttgtgtgtg tgtgtgtatg tgtgtcttgg agag 54

<210> 288
<211> 457
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 288

cccatcaact gccctaactc tttcagactg gtgattccta gtctttgac cttgacttga 60
tagaacctct tttaagcga aggccctga ctgcgttccaa tgtttacaa agtggaaacaa 120
aacccaatgt gaatcaaaaac tccgacatct atcatgggtg gaatggatga atgcttgaag 180
aatgcataat gacacagata catttatga atacgggagc ccggaaatt gtccccttct 240
tagatacaac attttggca gcatggcgcc tgacgtatgt atttaagaag gcgaaatgga 300
ccctccgtcg gtttgcacaaa gtgaggggac caagacacaa tccgtgcatt atgcataatgc 360
ggaaggcaca aaacggtgat gtacatagta cgacaatatc cacaacaaa tataagcaaa 420
ggcatacatg acatttanga ctacatgcat gacagtg 457

<210> 289
<211> 402
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 289

gctagagctt agtacacat acctctctaa tagctaagct caccccttg agatgagaag 60
ctagagctt gctacacacc cnctataata gctaagctca cccncatgac ananaaaaaca 120
tganaataca aaanaaaagtc cttactacaa agactactca naatgccccg aaatacaagg 180
ctaaaaccct atactattag aatggccaaa atacaaggcc caaacgaaga anaaacctat 240
tctaataat acaaagataa gcgggtcatg cttagccat gggctgaaa tctaccctaa 300
ggctcatgag aaccttangg cttcccttg atctctagcc caatctactt ggagtcttct 360
acccaatgcc cttgcaggat aggattgcac cacatgtcat ga 402

<210> 290
<211> 457
<212> DNA
<213> Glycine max
<400> 290

cccatcacat gtggtaactg gtggcggtcg ggcgttggcacaacaagt tttccacatc 60
cacaacgcgc gcataaaaccc accatccctt gttccccacc tccaaactgat ctcacgtact 120

cccacgtgc ccatatcctc ttttctctca acaccgggtc cccatcaatc ctcccaagct 180
ttcccaacat caaagtaaaa cgacattcaa acagcacaag ctatcacagc caagcaaaac 240
agagcaaagg cagaaaactc tgccaaaaca ccaaccatat cacagtttt ctcacttaaa 300
gactccaata acaattcctt cgttccggtt cattaaccgt tggatcgact cgaaaattgt 360
actggaagtc ttttagtacat aagcctacga tttgaccgtt gggatctact agcacacatc 420
cagaactcat tgtacattac tctctccaca accagcg 457

<210> 291
<211> 219
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 291

tattttagcc gatgctctt ctcggcgtca tgcattactt tctatgcttg aaacaaaatt 60
gattggtctt gaatgtttga aaagcatgtta tgaaaatgtat gaaactnttg gagaatctt 120
tagaaaattgt gagaaatttt cagagnatgg tttcttttaga catgaaggct ttctttcan 180
agaaaaacaaa ttgtgtgtgc ctaaatgttc tactagaaa 219

<210> 292
<211> 544
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 292

ngggacgtgg atgaattcct gcattgtccg cacccggta tcctntacag ccgaccggca 60
cgcatgcaac ctttatgaac tcacggngga naagcctcga actttgacac tcccgacgctg 120
tcacttatta gggtttgttt cgtggagcgt ttggcgacg atagagaggg cgtggaggcc 180
gctctggagc tcctcggcga ggaggtcgta gacgagacgg tgcgtttga cgaggctctg 240
gccctcgaac ttgggcgata cgatctcac gttgaagtgg gtttcttgc cgaaacttcc 300
cttcacggcg gcgtggcccg cgtgctggta cgacacgtcg tccacccca aaacgggtggc 360
ttccagcgcc gtttgaagct tcgagcgaat cctgctggct cgagatagca gcgcgcgtggc 420
tcctctcgaa ctcattggat cgatgaatg gtggcgctgg gttangttt tgagaacgg 480

ggagttccg cagatganaa catggacaga aattggggag ggaaagttat atttcccctc 540
tgcn 544

<210> 293
<211> 480
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 293

ntgaacccan agtaacaact atgacttggc tcaacaacac ttgttgcgct tactctcaac 60
cttcaaaagc aataactccc taattgatct tttaagattc cctatcctaa atgagtttg 120
tttggggaca aataccta ataaaaaaac tcaccaataa gtctctattt agaatatgtt 180
tacatacatt ctatggtaa gtgaatttgg ctctaaaggc ctatagagtt atatatctac 240
aaaatgactt ctccccc caacatagtc atgagttntg ctactttaga aaagtaagcc 300
ttctttcttc acgtaagtgc agttttttt tcttggtttc tagttgtatt gcttgtataa 360
gacttatgt gacaaattaa tctctcattt ccagaatgtat tcatgtctt tggagagctt 420
gtcacttcat atgcaaataat attgttgcattt ggcatttcct tccttgcttc taatgtgtgg 480

<210> 294
<211> 424
<212> DNA
<213> Glycine max
<400> 294

ggagttta catatatgac aaaggcgcaa cgcggtatgg ttgaaaatac cttctggc 60
tttactaaa aatataggta ttagatctgg agtacagata atcaagctaa aggtataaaa 120
gatagcctat gtggatcata acactataaa ggtgtgacca ggcttacag atttctactg 180
ttattatatt ctgtcttttgc ctctgactct gataatttac aagatcctt ttcataatgtc 240
tccgcaccgc ttcacattct aattcattta cgtgtatatt tctttacact ttagaaacta 300
catccatcaa ccatgccctt aacgtctaaa tctgtgaccc gtcatcgatc aagcagaagc 360
ggtccaacaa aagtgaaaag acacgatgctt gactttccca tattggagag atattcctca 420
gacg 424

<210> 295
<211> 353
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 295

cgacacataa ctcccaccgc atatagaata tcgggccttg tattggtagt ataccttata 60
ctccccacaa gactcttgaa gaccgtggaa tctaccttct ctccttcata aaactttgat 120
agcttcaagg caccttccat atgtgtgttc acgggattgc aatcaagcat attatata 180
ttcaaacattt cttttgttaa gctttttgtt gagacacaga taccattctc ccgttggta 240
cttnccattcc caagtatatg acatgagtcc atatttgcattt atcaattcac agacatgact 300
cttgagtcctc aacaatttgg tattgcgata aataggcatc cctaaacaat aat 353

<210> 296
<211> 436
<212> DNA
<213> Glycine max

<400> 296

ccagtttcct gaaaatatca aggaatctcg ttagatggcg gtcgttgtcc ttcttggaaag 60
gtaccacagg atatggtaact ttgttatcct catttgaagc ttttcttgc ttcttctctc 120
ttgcttctc acttctactc ttttcttcc cttctttatt ttttcaact ttttctttt 180
cttcatttcc ttttcttcc tctacctcta tttcttttc ttggcggtt atttctttct 240
tctcgaccgt tattggttt tcactctcct aacttgcac atctgtgcc tcctcttct 300
ttttctcaat gccatcctt acaacaatat gtttctcaa agccacccta tcctcatcct 360
caactaccaa acgcttattt cttgtcatca caacattaca ttccctctgtg ggattctctt 420
ctgtgttcgc cccaaag 436

<210> 297
<211> 365
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 297

ctctacaatt gcatcacctc tcaatgatct ggtgaagaag aatgtggcat ttacctgtgg 60
tgaaaaaaag gagcaagcct ttgcttgct caaagaaaag cttactaagg ctncagttct 120
agctcttcct gactnttcta anacttna gctagaatgt gatgcctctg gagtgggagt 180
tagagctgta ttngtacaag gtggcactc tattgcttat tttagtgaan aacttcata 240
tgccaccctc aactaccca cctatgataa agagctntat gccttaataa gagccctcca 300
nactgnnga atataccctn gttccanggg aattgcattc atagtatca tcaatcactt 360
aagta 365

<210> 298
<211> 453
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 298

aaggccaagt catcacgata tgcgaggatg actccccgag caagttggat ttggatgac 60
catgccctcc tggtttctga ctaggaaatt ggcgagtgga ggagcgccc aacatttacg 120
cgacaagcat aatgtAACCC tttgtggctt ttaaactcta cggtggggcc taggctttag 180
agtttccctt tgTTATGGCA ttatgtctt tgTTCTGAA tttataaata taaagatctt 240
tcttcatctg ttccTgcacc tctaccatt ctcattcatt tgcatttta tttcttacg 300
cttaanacac tagatccaac aacgagtccc tcgaaggtaC taatacctgn gaccggcca 360
tcgattcaag caagaagcgg gtcaaacaga gagtgaagag gacgaggatg tggacttcc 420
cccagagttg gagaagatag tcactcacga ggt 453

<210> 299.
<211> 479
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 299.

nggaatggca cttggaccac ctatTTGAAT ctccTATGCT gtacctacat acataaaaac 60
agtcccacca tctcaatttt tacaaaatca tattcataca ccattgggc atttcaccaa 120
gcacttggtg agcgcatgtt tggacatgaa ttgcaagaga atgggagcaa tgtggcatgc 180

cccattgctt cagaatacaa cctaggccta agacctttc attcaaatcc tcaattcaag 240
aaaacaagca ccaaagcaa ccaaactgc ctcacaaata taagcatgtt ctcacaattt 300
aaggcaccaa aagatgaaga aaacacatca atgggaagca aaaacatcaa ggatggaata 360
cttacttgtt ggagtgaatt gaaacaccaa aaatgaaagc aaaaggcaac caatagtggc 420
ttgagggggc aagaaccaca agccttcgtg ttcttcttt cttgaatgag aagggggan 479

<210> 300
<211> 541
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 300

cggcacatg tttgttgc tgcgtacccga gaccctctag agtcacactgc ggcacatgcaag 60
ctnngaacaga ttaaagtcaa caacactgac tggtnntggtt gagaggtgga gacctgagtc 120
acacatattt catcttctag ttggtaatg catagtcact ttggaaagatg ttgctcttca 180
cctgagttta tgcgttgatg gaaaaccaat taatgaccta acatattatg attgggaaca 240
aatgtgtgca aaatatatac gtgttgc cccaaagaat gcactagtgg gatcaacact 300
aaaactaana tggtaaaaag aaaacatgct gactctccca gcanaatcca cgcaacaata 360
attagcaccc cattgttaggc atacattnta ggaccaatta gacaagttag gaaacanagt 420
tcacctgatg tatctacctc tgtagcaaa tcttgcacag gcaggatggt acaattgaga 480
attgacatgt ttagcacatt tgtacagaga aatgtttatg acaatttatac catcatcaaa 540
n 541

<210> 301
<211> 221
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 301

gcctttcttc ttccctgaaca cacatggtca ttaattcatt gatagaccac ttatcttat 60
gtgtcggtta gaaatctta aatggcccat attcatgccg aagggtgttc agaatgaaat 120
gcactangaa ggactcagac atatcaacct ctatgtttt aagtngagct gaaatatctc 180

gcattntcat gatgtactca cgcatcacct tcacacttgt g

221

<210> 302

<211> 170

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 302

agcacaacta gtcctatgtc ttctcttga gagatanaga tactcataag agttgatgta 60

ctctactata catagttctc tcactgtgtt ggtcaacttg atgaactctc tcaagtgttt 120

cataggatct tcatgagcac caccactaaa tacgttgtct atagcatctg 170

<210> 303

<211> 256

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 303

gctcgaaaga gagttatgcn ctgtacactt anacagtgtt cagagtaata tattatgcc 60

nnaataagag aacatgacac ttggacccaa ttagtgc当地 tntcacaaaa ctcaattaan 120

aggcttcana accataataa aacatgtcan atatatgcaa aatgaaaacta taatgtatgc 180

tctaataattc tctatcagag gacattcgat aggaacanaa tgaagtccct tanacaatat 240

tcttatttgat gatgat 256

<210> 304

<211> 260

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 304

catgtggact atgtggcggt cgggcgatgg tgctcaacaa gttntccaca tccacaatgc 60

gcgcataaac ccaccatccc ctgatgccc cctccatctg agctcacgta ctaccacgta 120

gccccataatc ctcgttgtct caacaccggg tgcccatcaa tcctcgacca gctccacaac 180

atccaagcga aacaacattc aaacagcaca agctatcaca gccaaagcaa acagagcaca 240

ggcagaaaact ctgccaaaca 260

<210> 305
<211> 523
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 305

nccacggna agttccccag nnantcantn anatnganan naancaaana naataagaga 60
acaatgaaaa tggagaatt gattcatgtt tccttgatg agtctaattt tatttgtcca 120
agaaaggata ttttagatga tattgttagaa tccttagaac aaatgcacat tcatggacaa 180
gattctaaag gaaaaggaga aggaagcaat gaagatcctc cagtagaagt caaagaaaat 240
aatgatcttc caagagagtg gaaagcttca agagatcatt cccttgacaa cattattgg 300
aatatctcaa aagggataaac aactagacac tctctcaaag atttatgcaa taacatggtt 360
tttgtatcta taattgaacc taaaaattta aatgaagcca taatagatga aaattggata 420
atagctatgc aggaagaact ataaccaatt gaaagaaata atggtnntgga gttagttgag 480
aaacctgaaa actacccaat cattgaaaca aaatgggtgt tag 523

<210> 306
<211> 468
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 306

agactgagcg cttatcacaa ggtctgtgct tagcgatag acaattgcaa aaaaaatttc 60
taagtcttt tctgtctata tcacacaca agcttaaaac cccttgatca ttactaaaca 120
aactaaaaatt aatcacaatc acaatcaaga tatcctaact acatgcaaga ggtataatg 180
aaaatagaaa agggaaagaa aagcttagtt gcctcccagt aagcgcttt ttaacgtcac 240
tagcttgacg catcgctctg ttatccagga accaagagag ttccctacttc aaggaccttc 300
ttctcaggtc tctttcttc catcacatgc actntanaat aaacatntg gcttaggtgga 360
tccttggctc cctgaaacaa atcaaagctg atcttctgat cttctatgcc catccgcagt 420
atcttttttc ccatgtncac cacacagctt gcagtagaca tgaatgg 468

| | |
|-------|-------------|
| <210> | 307 |
| <211> | 472 |
| <212> | DNA |
| <213> | Glycine max |

<223> unsure at all n locations
<400> 307

| | |
|-------|-------------|
| <210> | 308 |
| <211> | 475 |
| <212> | DNA |
| <213> | Glycine max |

<223> unsure at all n locations
<400> 308

cgttgggtgc atcagtcggc agaccctatn atcaaactga ggcggcgtct cactgagagg 60
gcacgaccca gactgttctt attagaggtg gataccggac tcaccattac tataatctct 120
accctgtgca tgctctgggg ctccagtaga tgccggctcat cccctgagct tatacgcccta 180
tggtataccca gatacggacc cttagatgc aagatgtctaa caaagcgtgc gatgcggata 240
tacatgtaga tcgcccgtat actgcctaat gggagcacgt tctggcccta tggataggaa 300
agcgtgtatgc cagctagtga cctcacgcac tttaacttagta gactttgtgg catgcatttt 360
agaccatcta cagtaggaac cccgcacgtg ggacctcctc tggacatact ggacagcagc 420
ggtcgacttag attgcacgcc atccactgca gagaaggtat aaattatcat atcaq 475

| | |
|-------|-------------|
| <210> | 309 |
| <211> | 511 |
| <212> | DNA |
| <213> | Glycine max |

<223> unsure at all n locations
<400> 309

gtgatgatgc atcattatct acgngaacat actcaagctt gctcgacg aggtactac 60
ccgttgaaga tcgatgaacg ttgattatcg aatgaagaac gttgaagaac gggtgatacc 120
tttgagagat tcctcaccga caacgtgcg gatacgcatt cgaatgcgt ggccttagat 180
tgacttgatg tggacaagta atgcgagcaa attggattga cacataggta cctaattggc 240
tcaacgcctt aattcttgc tttctaacct atatataaca aaacaaggaa cgtggttgac 300
gcccgactcg cccaggcgag ctcaactcgc ccaggcgagc agggttgc ttccagaag 360
caaccgcctt ctggaggaat attccggagg gcccaagtgt gcctgggtgc tatttgcacc 420
cccatcttta ctaagaacac cacgctacgc tgtttcggg gagggtctat aatacagtac 480
cgtaacttac gatcgctga agaaaggggg g . 511

<210> 310
<211> 421
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 310

agttgcacat ggattttct canaacattc ttaccaaaga gttttactc tttggtaatc 60
gattaccaga ttattgtat cgattaccag tagcaaaatg gattgaaaa agtttcaaa 120
ttgaatttac aatgttccaa ttaattcaa aaagctgtaa tcgattacaa tgtttggta 180
atcgattacc agttccttg aacgttgaca ttcaaattca aatgtgaaga gtcacatcct 240
ttcacataaa agcctgtgt aatcgattac actgattgg taatcgatta tcaatgatta 300
tttctgaata aatcaaaaga tgtaactctt catattgtt tgattttca catggattaa 360
gctctctaaa actatactct tcataatggc gatngaccaa cttaatagtc atacatttc 420
t 421

<210> 311
<211> 234
<212> DNA
<213> Glycine max

<223> unsure at all n locations

<400> 311

acatcatcta ctattgttca tctgcttcca tgaatgaaga ttcatgatca tcacaggtac 60
cacaccacag gtacaanaat tgcagggtga gtntattata aaagaaataa tcaaggcatta 120
gatgacaata attagcaagg aaactataac aataaccata atcatactta ataattcatc 180
agtttgacat acactanaca tctagtcatc aactttcatc atttncaatc aatc 234

<210> 312

<211> 221

<212> DNA

<213> Glycine max

<400> 312

aacagttcaa tcacatgccc ataaccacat cctgtgcccc tcactgagct agactcacga 60
ccattctgtt ttaaacggtc catatctcca agttccacat aaggatcaca taccagccag 120
tatacagcag caacaggcaa gaaaacttgt caaaaccaca aataagttt actaagaaca 180
gtacattctc atcaatcgta ccgtgatcga tcaaattatg a 221

<210> 313

<211> 455

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 313

ctgagcaaat tcaaacgaca ataactntat aatcgatgt cctattgagt cccctaata 60
atcaaactgc tccaaattga aaatgaaagc tcttagcata tttaaacgag aataactctt 120
tactcaaatg tgcgatttag tcacgtaata tatcgagacg ctctaaattg aaaacggaag 180
ctcatagcaa atgtaaaccg taataacttt taactcgat gtccgaatga gtcctgtat 240
atattgagac gctcataatt gaaaacagat gctctgcgc tattctaaca acaataacct 300
tttactctgt tgcgaaatg agtactggaa tattngaga ccctcgaatt gaacacaaag 360
ctcctaaaaa atcaaacaaa aacttttatt ttatgttac tgaaccgtat ttccggacgc 420
tcacatggaa caaacttctt tattcaacgc agtcg 455

<210> 314

<211> 446

<212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 314

 tcaaggaagc ttcttaagga agtttctcaa ggaagctacc tagtctataa atagaagcat 60
 gtgttaacact tggtaact ttgatgaata agagtcttg gagacacaac tcaaagttca 120
 acttctctcc ccttttcct cttcaattt tgtgctcccc cctctctttt tcttttcctc 180
 cattgaagaa tcctctccaa gtttcttac caaggcacat tcttggtgc gaagctcctt 240
 ctccatggc ntntcccta gaggatggcg cctttctcc tttgtcttcc actgcatttc 300
 cgtggggaa aatcaccatt gaaggaccc attgaagctc aaagatccag cctncataga 360
 agttcacaag taagttcat catattctct tangcacaac actgtggcag tatggactac 420
 cagcgacaat gcatcaccat naaaat 446

 <210> 315
 <211> 482
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 315

 ctgatggtgt cgagaagaga tcacatgttt gtcacatcatca aaaaggggaa gaatgtgaat 60
 gtatgtatac atgatggta tgatgtcaaa gaagaatcta acaaggctac ttcaaatgt 120
 aacatttgc ttcaagaata attcaagatt gttcaacaa acaaaggctt gttcaagat 180
 tcactaaaga ccaaggcctt cttataaca aagtgcattt aagacatgca aggctctgg 240
 aatcgattac caggaagtgt aatcgattac ccgaaggcagg gttgagaaat agctgttgaa 300
 aaaggtttg aatttgaatt ttcaacatgt aatcgattac catatgtctg taatcgatta 360
 ccagcaacga aactntggaa attcaaattt aaaaagtcata acccttcaaa ttataactgt 420
 gtagtcgatt acacacacat tgtaatcgat taccagtgg aagttcaga aaatctgcc 480
 cg 482

 <210> 316
 <211> 441
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
<400> 316

attaggttgt cagttctgca aagataacaa tgcagacgta gcataattac attatagtaa 60
aaaaatttgtt ttgcagacac ataccttgat tttaaatttt tgaacataaa cgatcataag 120
ctctagaaaa aataactcacc agtctccaaa aacatggttg gcaatggcac atggtgagtg 180
gccttggcct ccaatgtaat ctaaattagg gaaataaaatc aaattctacc aaaagtgcct 240
ttcaaatttc aaaatgtaga cttaaaaaac acaaattaac actatgtcat catcctccaa 300
caaaaaacaaa cctgtgtttg cattctatga agtgaactat ttgtttaaat gattatgcat 360
gaagacatta nacacagttt taccataaa tatgcaaact atgagaatnt canacaactc 420
anataccaag ccaactatTTT a 441

<210> 317
<211> 262
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 317

cattctagca gntccttatg atataagcta agtcaatgac cagccttang tttcgtagt 60
aggtgagagc atcagatcca acttcccttg atctacacaa ggatgtgatt aaagctggga 120
agcctanacg agaagagttt gatngagcca tcatggtcat ttgtcttagag atcaaaccgc 180
caatgttcat gtccatcctt gtgattaagc catagaccga cctagctcta tcntgtgtca 240
natctgaaat gaaggatgta tg 262

<210> 318
<211> 537
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 318

ggtgactcca tncnnnnatg aaacctctcg tagtaccgt gatcctctag agacgacccc 60
gccgcattgca agctntgcgg atttggtctt cgccagtgaa atgatcgaag tggatctgaa 120
aagaggcaaa tttaatcatc ctgcttagac gaatgagaaa actgnggcaa ataaagaggg 180

tgaggatgag ggagaaaccc atgctgtac tgccattcct atacggccaa gtttcccacc 240
aaacccaaca atgtcattac tcagtcaata acaaaccacc tccttaccca ccacccagtt 300
atccacaaag gccatcccta aatcaaccac aaagcctgtc taccgcactt ccaatgacga 360
agaccaccc tt tagcaaanan caaanaaaaa caccaaccaa gatatgaatt ttgcagcgaa 420
nagcctgttag gattcacccc aaattccgggt gtcatatgct aacttgcgtcc catatctact 480
tgataacgca atggtagcca taaccctgc tagggttcct caaacaccc tc atttttg 537

<210> 319
<211> 512
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 319

nggcgtgcta tgtcctgagn atcnacngat ntgaaaaacc aagctttatt ttgcttgatt 60
ctctgcaatt ctctgcattcc ttgctccctcg aattgaacct tcaacctttg acattatttt 120
ctccttcttc actatgaggg aaggtaactt ctttgtgtga gtgctttgg ttttggtttt 180
gacgtaagta ggaggaaggt tataggtaaa aaaaaaatta taattatttt ttaatcattc 240
tgtagagaag aagttgttta ttaagagagt gggtagaga ggttaattaa taatggagaa 300
gttataactc ctcctttat ctaattgatt cacatgatat tttagaacaa acacaatatt 360
tccatttgta ataaacattc ataaaattaa attcttcgat taattagcat gaatggtagc 420
agtatgaaga attattattt ntatcaatga ggcaagagat agttctaatt ttataaaaca 480
agttaaatat gctnttagtc cataataat tg 512

<210> 320
<211> 533
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 320

tggacaagtt gacgcttgta aactggacca tagagnacc ctgcggcatg caagcttattc 60
gctgacttat tatccacaaa aagcttcact ccatcactct cttgatnt taattcctat 120
aataatgtgt ccaaccagac agcttggcaa gcactcattg cagctgcaac atactcagct 180

tcacatgatg ataaagccac tatggattgc ttcttagaac tccatgatat tggtgttgc 240
ccatacatga atatgtaacc tgtagtactc ttttgcatt ctctgtctcc tccccgatcc 300
gcatcagttat atcccactaa ttcttctgag ttgggttgttgt ctttatttgg aaatagaatt 360
ccagtattga tggttccttt tatgaacctt agaattctct tagcagctag gagatgagga 420
attctgggtc tttcgtaata tctacttacc agtccaacaa caaattccaa atcaggtctt 480
gaatgataca agtacctgat gagagaacca acaatctggt tgaactcagt ttn 533

<210> 321
<211> 301
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 321

agagacatga aagaccggat gagnttact gtgagatgga agatctaact tataagcaac 60
aacaccacc ttatntaaca cctggaaagg accataaaat cgangggaga gnttttcatt 120
aatccttna gccaggatc ttctcttgta ggggcattct tcaagaacac ccaatcaccc 180
actgtgtatt ctatgtcctg acgaacgttt gtggcatng ctgcattgtat atcttgagac 240
ttcaacanga tttctcttat agtagccaat aattaatcta agcaanntt tagttattga 300
C 301

<210> 322
<211> 450
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 322

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actcattcaa attccacagt ttccctact caaatcccc agtaacattc tcttagttcc 120
gattcgtaa ccattggatc accttggaaac gtttactgga gttccctagt acataaatct 180
acattttgcac cggtggatc tactagaaaa tatctagaac acgagatata ctaccttcc 240
cgtgactggt gctgcacaag catttttct gcacatttgg tcaagttgc tgccacaattt 300
gacagctttt gctgcacaat ttggcagatt tcanaatcca actttccac antccaattt 360

| | |
|--|-----|
| actcannatg gatcctanaa ttcctaaatc atgtataaaat catantaaa ccanaaaca | 420 |
| acttcagacc aaggcanatc anaatatacg | 450 |
| | |
| <210> 323 | |
| <211> 362 | |
| <212> DNA | |
| <213> Glycine max | |
| | |
| <223> unsure at all n locations | |
| <400> 323 | |
| | |
| ctcagtatgt tccttggta gataatata aattaactta gatacaattc acaattgtat | 60 |
| ataancatta natatgttat aattaaagaa ataactacct ctcttgccta cactttggct | 120 |
| accacatgat taacatatga tgtcaacact aatgtatctt gggcccccacc tgaaaaaccc | 180 |
| tatgaatcaa cacctacatc ctttgtaatt ggatcatgan ngttctcatg agtctcatca | 240 |
| gcagcatcat cgatatgccc attatcctcg acaataggtg cagttgtca ttatctacgt | 300 |
| gtcgacactt tcaatcttcg acgctgangg gcttcttctc gtcaccacta acctctctac | 360 |
| ct | 362 |
| | |
| <210> 324 | |
| <211> 531 | |
| <212> DNA | |
| <213> Glycine max | |
| | |
| <223> unsure at all n locations | |
| <400> 324 | |
| | |
| tgatgaatca tctcganctt gagatcctta gagacaccct gcggcatgca agctntgacc | 60 |
| gcatcttaac aatatctttt gttctattnt tgtgttgttn ttaatcaatg tctttaagtg | 120 |
| catcttaaca atgtctttt gataggatg aaaaaataga cactaatttt gtgggtatct | 180 |
| ataaaagtat ttgcaaatacg gaaggataat tatctgctta ttggactag agatggggcg | 240 |
| ngnatactat agtaccatct caccctccc cgcacatgta tgtcatatat ttatatatatt | 300 |
| aatgtatcca aaaaataatt ataatttcct aatttatga ctagcaataa caatctaaca | 360 |
| aagaataaaag aatcctaatt caacattgtt atattaaatt tgcttcacat tctacataag | 420 |
| aatatcgaac tactatttac tgtttatgtt taaagtaatg actctgctaa caagttatta | 480 |
| taagattgac ttgtgaaatg gtattctgtg ttgttcgatg cttcaaacta g | 531 |

<210> 325
<211> 470
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 325

atttcaatgc ggaaagtatt atgttcttca ctatccatgt tcacacatta ttgctgcttg 60
tggttacgtg aacatgaatt acttccaata ttagatgtt gtttacacaa atgagcacat 120
cctanaagct tattccgcgc aatggggcc tcttngaaat gaagcgagta ttcctccttc 180
tgatgagcaa tggacactta tccctgatcc aagtacaatt cgtgcgaaag gtcggccaaa 240
atcaacaagg ataatgaatg agatggattt gctggaccat ctgacaccga caanatgtnt 300
agatgtgaag agaagaccac agacgtgatg tcaatgaatc tgatgtggaa gttgtaataa 360
tgattatgtt tttgttgtca cttaatgaat gacctatcat gacagctgtt ttaaatagta 420
tatatatattat ggcggcctaa ctgacaatgg taatataac ataatgatat 470

<210> 326
<211> 315
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 326

ctgcagctag ctgctggcag ctgggtggaaag ctcccttctt atttnnncta taatagggga 60
ggagtgttggaaagg agagaaatgt tcagacccatc tggatattcg agatcaacttggaaatattgtga 120
aaaanactgt ctccgtgaag aaaatacaag ccgacgcgc ttcgtaacgt ttcgtggggaa 180
tttcgcgaag aatttaccta tntcttcgac gtcttcgttc gttcttcgtt cttcaaccccg 240
gtaagttctc gaaatcgaaa cttcaatttcc attctatgtt cccttagtgg tcctcatttgc 300
tttcacgtgc tttat 315

<210> 327
<211> 352
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 327

ctctctctnn tcgaattgtt gaggaagatt attccgtga agaanatcca agccgagggc 60
gcttcgtaac gttccgtga gtaattacgc gaatattctc gaccgttctt canagattca 120
tcgttcggttc ttctttctc tcagtcttca acggtaagt acctcanaca aagctttca 180
attcactcta tgtacccgtg gnggtccaca ttntggttca tggagttta ttctcattnt 240
catttactnt ntataacccc ttttgacgtg cttaaagccat ntatttaagt catttctcgc 300
ttaatctacn aaataaaaata aatttccacc gatcggttga atcgcatatc cg 352

<210> 328
<211> 514
<212> DNA
<213> Glycine max

<400> 328

ctttgactgc tgcatctgct acccttagag acgaatcgag gcatgcaagc ttagcaccac 60
tatcgcgctt agcgcgagta aatggatttg gacttggcgc caatgttgca ctgagcctag 120
caagagatgg acgactcgct tagcgagctg atctcgcgct tagggcgctg cttcgattca 180
ggtgctcttc cagattcctt tttcacgcta agtgcactga agccgtgctt agtgacggat 240
acgcactaag tccactgagt tcgcttagtg cgacacccag cttccgactt gaagacatca 300
gtaacttatt atcttagctc ggccaaagtc tacctctcct catctcacag aggccccacg 360
catcgacta gcaccgctgc ctgttattcg tacaagttagc tgacaactat acacaggtac 420
ccttcatcta tcgcatctac ctcaacgcaa gcatcagcta ctgtacgtat cgtctcatct 480
ccgcacccctc tcgacgactg ccacgagacg tccg 514

<210> 329
<211> 341
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 329

tctacacctg ttgcaagagt ctgtggtcta tggttcttca cagatcacca tacagatctc 60
ggcccttctt ttagcaatc tggaggtaat gagcaacctg aagcttatgc tgcatacatt 120
tataatagac ctccctcagca gcaaaaccaa caacagaaaa ataattatga cctttcaagc 180

aatagataca atctaggttg gaggaatcat ccaaattctga gatggacaag tccttcacaa 240
caacaacagc ttatcgcttc tttctagaat gctgctggtc caagcaagcc atatgtntct 300
tctncaatac agcaacaaca gtcacaaana agacaacaag c 341

<210> 330
<211> 450
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 330

atgtatccaa aaccctcaa tttaatggat tttcaagttt tgagaagtga aattggaaat 60
gggataaaatt tggagcaaac tctcacctca cacaagtcta taacatcaat ttaaacttga 120
tcaaactgga tttacaccta aaatttcaact gaatcagaat gtgactactc aacccccaaat 180
tttaccctag aaatggctct ttgttcagtt aggtcatttgc tttttcttt tagcacagcc 240
cagactttct cataagtccct atatgacatt gcaagctagg attaactgac tgtaacctcc 300
acataaccact aaatccagat ttggccttcc aactatanaa cattcaactct ttttacactc 360
ataacaccat aatctcacct tctaaccctt ggttaattct acacttcatc tctaacagat 420
ctccataagc aagttcagca cacatacatn 450

<210> 331
<211> 471
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 331

cgtcagagtt tanggattga gcttggttca actgagtgtc catctgcccc atctgattgg 60
tcaaactctg aatggaggct cttgtctctt gctgaaactg catattttgtt attgtcattt 120
gccttactaa ctccctctaag gaagggttag aaggggcctc agttgattgt tgcctctgtt 180
gttggcgtg ttgatgctat tggatgtttt gcattggagg aggaatgtat ggccctgcttg 240
gaccagcaac attttggaaag gaaggagcag gttgttggat ctgttggat gggctagacc 300
atatgagatt agggtgattc ctccatccga gattgcattt gttgctagag aggtcataat 360
tggatgttgcata tggatgttgcattc tggatgttgcattt gttgctagag aggtcataat 420

catatagctt angctgctca attgctccag attgctgcat agaanggcac g 471

<210> 332
<211> 446
<212> DNA
<213> Glycine max

<400> 332

gtttccgttg ttcaatttcg agcgttaga ttagttatgt ccccgaatcg gacatctgtg 60
tgaaaagtta tgaccattcg attttctcga gagttccgt tggcaattt cgagcgtctc 120
gatatattat gaccccaat cggacatctg tgtgaaaacg tatgaccatt cgattttctc 180
gagagcttcc gttgatcaat ttgcagcgtc tagatgagtt atgtccccga atcgaacatt 240
cgagtaaaaa cttatgacca ttcaatttc tcgagagctt ccgttggca atttcgagcg 300
tctcgatata taatgtcccc gaatcggaca tccgagcgaa atgttatgac cattcgatct 360
tctcgagagc ttccgttgc aatttcgagc gtctcgatat attatgtccg cgactcggac 420
atccgtgtga aaacttatga ccattg 446

<210> 333
<211> 400
<212> DNA
<213> Glycine max

<400> 333

gagtgattca agaacaccct gtctgtatca tatgacattc acaacctttg cgtgttgc 60
tcgctggaaa gagcgagtct ttccctcctt tcatactatac ccgttgcatt ttcaaaccac 120
aagtccagaa gatacacccct tgcccaagaat tatatcgtgg ccataactcc cattctacgc 180
actcacatta agtgattctt gagcctatac tgaatttcac aacgagttct ttccaccc 240
tatggaacac ctcattggag ctttagctca gtatgtcatt ctaattttgt caccacactt 300
actagttac atccattatc atttatgcaa gaccactt agacacgaat acactattca 360
ccttctataa tccctttcat agttatcaac atctagcact 400

<210> 334
<211> 457
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 334

tatacaatgt tntcatgata aaaggctctg ttgaccatcg gaacaagatt tccaaagggt 60
gcctctggaa atgcagtccct tgcattcatg aacctgaact gtgagatgaa gatcgaagat 120
caaagatcgg atggttcatg tctcgaatac ttaatttaaa ttagattggg ggagataaaa 180
tataatttgtt ctgatcttca tacaatggtt gcgatttagcc aaatgcctga atatggaaaa 240
attccctgac tgccgtgaat cccaatccct ctaattaaat aagctttaga atgcagcact 300
gaactaaaaa atttcttgcg tcatttatatacaatcttagt aagtaatgca ccaaagttcc 360
aattttgca tcanaggaca gagctgatag cacataaact aaatggcata aaaatacaac 420
caaatcttat cactgagttc tatcaatgga ggaggcg 457

<210> 335
<211> 260
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 335

ctatcttaaa gtaagctcta tgaaatttat aaatcgatat ttatatttct aacacgatct 60
tccttacaac tgataaagac aaagagaaca aattcaatta taaaataaag gataaaggaa 120
aaaactcttg acccacacac acaaaaagaga cgtattatag taaaaaatat ttaaggcatg 180
aaggataagc aagagtggnt ntaatttata attctgacgt tctccttgta attcccattc 240
ataattttct tctgtacttg 260

<210> 336
<211> 446
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 336

tccttcaaataacttgcaaa ttatggttct agaattctat aattttctat aggtttcagt 60
ctgcttagcta gttatttagac tggatggattc tgaatagggt ttatccaga tttattgtat 120
tttttggtttc taggtaatgt ccagtcatgc ttctttaaaa tctaattatt taaaaccaca 180
aaatttagag cctgcacttt ttaatgtgtt tatgaaattc ttgcgtttta gctattcatt 240

atccttgcaa tctcttattt cacatggtag tacaattcan attaacttca gattttgggt 300
ttatttgat gttgctctc gaagcagtct ggaattttcg gtcctttag acgggttcgg 360
ntaatccctt cttatnggtn tgtctgtact ttgttaggaag cactcaagac tagtgtggac 420
gcttcagtcg ttcccacatt tgagag 446

<210> 337
<211> 255
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 337

gatcttgct tgcaactgct ntaagtagtt aatcatctca tctaaccattg aagctntatc 60
tgtctgcaat tttgccaagt aatccaccac gtacaggaat aatcaatcca catatataga 120
acnntgattt tgcatngat tntggatnga atatgnntnt aactatgtaa ttnttgataaa 180
aaatgaaatg atattagtct ccgattctaa cttaatcaa tttaatccctt annatttaaa 240
agactgttnt cgtca 255

<210> 338
<211> 441
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 338

ganaactacc aaaactaccc atcatatctc ccaaaacccc atacccacga aatttaagag 60
agaaaagaagt ccacccaaac ctgaaatttc gaagtccac tcgtagccac gcacttcacg 120
actccaaaaa tgccctcctt tcgcgatttg gagcagaaat gagcaccaaa gggtgaagct 180
ttgtttggag cttcaatgga gaatgaggga gaaagaaagg caacgtgagg aagagagaga 240
gctgtctgaa aaaagtgtgg gggctgagtg aagagagaga anagcttnt gggtataaaa 300
taaaagggtt ttctctttt ctattatnt attcanactc tgccacgtgt ccctaattga 360
gtggagcana agggccccact ttctctttt ctgtgaccca cactcagcca canaagttag 420
aanaatctga ccttganac t 441

<210> 339
<211> 535
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 339

agagccgata ggtgggngc ttgnacnancgacantata tagtactcat tgctgccaag 60
aagggtggtgg atctcgaggc ccgactgaat gaattatagt ccatgctcaa ggagtctgag 120
ctacgggctg ctagagagag ggaggccagc aaggagctt aggaggagtt tttcattttc 180
aagaaggagg ccgtggagca gcatgaaaaa gggcctaacc aagccgttgg gcaggctggg 240
ttcttcacca aggaccttga cttgggtctc tttgaccctt ttaaggacgt gaagaatgg 300
gttttgcttgc acaaaagacga tattgctgct aaagaggagt aaggcgatga tgccattgg 360
tagggtgcct ttcgtttatt ttcttcttt ctccattgtt tgaatttagc cgcatgggcc 420
ttgttaattat gacaaattat cttcataaagc ttcccttcga tgacaaattn tgcaactatnt 480
atgtatgtct tgggtgttgg ctttatctat gcaatgctcc atgcttgttgg tagtn 535

<210> 340
<211> 406
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 340

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tacctgtgtt ngtgcagacc ttacaatcac tgccacanat ccattttcat gagcaacgga 120
tcgagccac cgcanaacat nctctggct gtcaataacct acaaccaatc agataatatc 180
agttctacaa catattatattt atttataaat actccaacat aacgtattac taaaagattg 240
acgcataac aacaatggat tatcacccaa tttctatattt ataatcaatc acttttaaca 300
taacttatca tcatactcg catacata ataaaattaa ccataaacac ccatcaacta 360
tctacatcaa cttctataact ctctgttaaca gttataaaca ataata 406

<210> 341
<211> 466
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 341

ttctacagaa tccgtanagt ttctgaggt ttccgaagga aaacaaccaa acaacacaaa 60
aattgggggg ggggtgaac ttatcaagct aggagtgtaa atagcaattt tcaaatcttg 120
gcagaggatt ctggaccttt tctttttcc tggctaagca acttggcga gcagggtggc 180
aagcacctcc ctcattttgt tgaaaaatgg ctccggcgc ttccgtagaa ttcccgtAAC 240
cataaataag tatatttcac ttaatatgg tgagaaggaa gaaaaaaaaaa gaagaaaaatc 300
aagtccgata tgcttccgta acttttcgt aaattacgaa gaaggggggt gaacttatca 360
agtgcgaggt gtaaatagaa attttgaac tttcgaatct cggcccttcc agaacattat 420
ggaagctcggttgcgtttagg agggagcagc ctacctcgct tgggcc 466

<210> 342
<211> 444
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 342

cataacattt ctaagtgata aagttttctt ctgtacttct ctgggggttc tgctcatgg 60
ttctttaca agtctagttc taggctgata tgcaagtggag ttggaaatct ttattccagt 120
aatgaccccttggatttttt ctgtttctt tcttactttt gttgtgagga tatcttattt 180
tcctctatac tggacacact tcctttctt ctccattaaa ctgtttctt tatagagatc 240
tgaacttgcgtt gatattgtgt ccaggttttgc caacagttgc agtctatgac tctgtatgacc 300
aatctcataa ttttcttgcgtt gtttcttctt tggtatttctt caattacaca tcgcgttgc 360
atactctgtt gatTTAATC cttctatac tagttgtca ttgtgttttttatacttctt 420
tctatTTAA tctcatgcag ctgg 444

<210> 343
<211> 488
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 343

atctccttct tcactacatc aagaatcacc gngttgagtc ttctctgtgg ctgtcttact 60
gggttagctc catcttctan atntattcga tgcatacatg tggatggct aataccagga 120
atgtccgccca gggccagcc tatagccttc ttatgcttct tgagaactga caacaacttc 180
tcctcttgct catcagcaag ggaggcagat ataatcactg gagaactctt gctatcatcc 240
aagtaagcgt attntaaatn tcatggcaga ggcttcaatt ctggtgtggt cggctggaca 300
gtggtagaag gagatggttt ctcagcctt acctcataaa gaaagtcaaga ggtatgtgta 360
cttcctgaaa catggtagt cctatctgac tctatnaaat caatctcaag aggtanaaca 420
ccaccaccag gcatgcantc aatatcactc tcagaatcac tctcagcatc anattcagac 480
atatgatc 488

<210> 344
<211> 532
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 344

ntttaatccc tcagtccccg agctccttaa gaaacccac gcatgcaagc ttcacggta 60
atctacagtg attcanagat gttntgatga taacaatgat gataacaaaa gatgatgaca 120
aagggtgatga caaaaagctc aaaggtaat taaaggatga gttcaagatg ttcaagaagg 180
aatcaagaac aattcaagac tcaagacgaa aaggttgaag gacactcaa gattcaagag 240
gaaagttgaa ttcaaaaatc aagattcaag gatcaagatt taagaatcaa gatcaagatt 300
caagattcaa ggttcaagaa ctcaagagaa gacttaatca agataagtat ganaaggttt 360
tttcaaaaac ttagtagcac atggatttt cacanaacat gtttagcana gagttntac 420
tctctggtaa tcgattacca gattgctgta atcgattact agtagcaaaa tttttttgaa 480
gttntcanat tgaatntaca acgttccatt taattcana aagctgtaat cg 532

<210> 345
<211> 290
<212> DNA
<213> Glycine max

<400> 345

catttgcgtg cttatccctg tatggatgatgatgaaatgc aaaggtaacg acttggtaa 60

```
gtcgggtata atggaatgag cctaaacact tgagctttag tgaaacgacg actgtgaggc 120  
tgtggtttag gatccttcct tgatatctgt cattctcact agcttatttc aattatgact 180  
ctaattgcata tctttctatc tttgaaaagt tgcatgtatg tgagaagcaa ttgattgaag 240  
cattccatga tattcatttc atatgattga atttttctgt aaacaaacac 290
```

| | |
|-------|-------------|
| <210> | 346 |
| <211> | 377 |
| <212> | DNA |
| <213> | Glycine max |

<223> unsure at all n locations
<400> 346

tatcacatat atatatatgt tatgcgtaca acatatatca ttacgcaatg acatttgagt 60
ataataaaaa atagttctgc agggcctaac atttcagtgc ttatattaat ttaggtacca 120
cttaacattt attattgagt caactctcta acgnatattc ataatttctc tttgtaatat 180
taattnaattt ggntaaagaa acatatttct tatggataat aatggcttcc agnttcttag 240
tgaaccacat ctgannaata tacttgacaa gaaatgtgtt actatgtcat agntaatctt 300
tttttctttt aaatacatca tctcttatgg acgattttag actcgagga cttattatat 360
ggacatacac ttatata 377

| | |
|-------|-------------|
| <210> | 347 |
| <211> | 396 |
| <212> | DNA |
| <213> | Glycine max |

<223> unsure at all n locations
<400> 347

| | | | | | | |
|------------|--------------|------------|------------|-------------|-------------|-----|
| cacacatgt | gtgaacatct | atgataacct | gcttcgtgt | agcattgtgt | tagctataat | 60 |
| ntatgaagaa | ccacttctag | ttctataatt | gtacaacata | ttagcatatg | ccanactatg | 120 |
| tgtatcattt | ggatcaccag | aataagaata | ttacctcaat | aaaatctcct | tttggcatta | 180 |
| gtgctctgca | tgcacatctcta | ntccttttgt | atggtgat | taaacttagtg | atgcaaataa | 240 |
| caccagcatc | tgcaaagaag | ttagccacct | cacctgaana | tntaaatcg | gatgtctaann | 300 |
| tattaataaa | acaataaaat | ataatcgaa | gatatcagg | anagcattta | gaaagcaaca | 360 |
| taagaaaaaa | cagaataact | caccaatcct | tctaat | | | 396 |

<210> 348
<211> 225
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 348

tagttacctt cttatgccta gccctatana tactcanaaa ctcttaattn taggagaatt 60
ttgtagaatt gaaattaagn tgtgcttaga gagagcatta gcctcttctt tggtnntgac 120
tagaaaccaa atggattctt ctcaaagaag ctattcctt atggcaaatc ctcctactcg 180
gtatcgattc ttcatggatt gtggcatcgn tctgtcatct tctca 225

<210> 349
<211> 203
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 349

acgttatcta tagaacattt ccattggatg taatggatga aattgtgcat ctttaggtga 60
gaaagaggct atgtttgaa ttgcanaatg tagcagttgg gctaaacgca tatccaccgg 120
taagcgcaat ttcaagcgtgc ttagtgcana ggagaatctg ggagagcatc aacatcaaag 180
ccgcgcgcta agagtggat tag 203

<210> 350
<211> 455
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 350

gacctaaatg aagactaaac atgcattgtt tatgtatttgcattat gcgatataat 60
ttgttgtaag ccattaataa ccaattaata ttattaagta ctcgtttgt taagcaaaaa 120
aattgttgtt ccaacaaaaa tcatttacgc gtgttaggata catcattgtc ataattgaca 180
acacataatg acatgcattgc gtattaaagt ttgagcgcga caccacatttgc actaacttga 240
ctacacattc tgaaggaaac ataaacacga aacatgttca tgcgtgtcta attttttgtta 300

aacaaagaga agcaatctgt ctgtgacaac catgtatata tatagcagac acaactaata 360
aatcacacat tatcttgctn tcacatagtc tcccaatgga tacacanagt atgaaatttg 420
tagagaaaact agcagtcaga tgattgcaac tcacg 455

<210> 351
<211> 483
<212> DNA
<213> Glycine max

<400> 351

atgttagtct gctcacatca aagagatata ttgtttctc tctcagatat atttgatcct 60
aatcttatcg ttttcttata tgcgaactca tcagctgtaa cattcttatac ttatctacac 120
acttgaggta atatttatag taattaatac aacttataatc ttatcttga ttaacctgtg 180
cagattgtta agcttcgag aattaaaaaa agaatttagac tcttgaggat cctgaattaa 240
acgtgaacaa gtatatatag gagcaagttt atccattgtatcattatagat tagaaataaa 300
aactacttgt ttgtgataga aataaatatt tatagttaac ccacaatgaa ttccgaaata 360
ttattattga taatttata gtgcaaaggat attacatta tatacactta ggctcataat 420
tgtttgccta gaataaaagca acaaactatt ccaactataa agggaaataa gtcagaaata 480
aat 483

<210> 352
<211> 448
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 352

tagcattaaa tgcacatcct tcttgatgca acaaaattac tctgattggc ttttgttag 60
aataacttcag tataaaacca ctttccttg gtcaaaggat gttttcaca tctgaatgcg 120
tctttgatg ttgttttaac aatttttaa ctttagtttc atttattatt cataggattc 180
gacaaatcat atgagaatgt ctctccaaca tgaatctcag acacagaaaa ataaatata 240
agcgaaatat catttttaa tggtgtatca ggtcatgact tggtcctatac ttcatctaat 300
acttttgcg catgatgtat acaacatgtat cttatcgataa actattcacc 360
ctcgtattta tgtgcacatcga ataagaacaa ttatgagttatc tatgacataa 420

atgtcncattat cttAACATAG atgaaatg

448

<210> 353
<211> 277
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 353

cctggagata tgcgtcgccc gtcaggagaa ccttgggacg tcaagtggng tgctatngcc 60
canaaccaag cttgaccaat cccgacccaa cccgcgcata gtcggtcagt gagaacctgt 120
gatgtaccta agcaggcgag ctccctggcag tcaacagata anagaaaaac aagaccacan 180
agcaaggagg cttgtggtgg ctggccagct gtgaattttg tgtaatatgt ggattgtggc 240
ctctggtaat cgattaccaa gggtgagtaa tcgatta 277

<210> 354
<211> 302
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 354

cggtgataact ggtttctaga agtggctcca gccattccca tgnnttcaca tttaaagaaa 60
agggagaaaac aaccaacctt atgtactcct tgtaatgtt gggtnntta acaatgtcac 120
acaacctcaa gaatctcctc anatgcatca caagaagctc tgnntgaact ggtactntta 180
atcatctgga ggaacccatg tgtgaactca naagtaactc ctcgaggaag tggtagt 240
tcaatgggtg tggttctcc caccagatgg gtgacataat acagcaatgt cagtgttagca 300
tc 302

<210> 355
<211> 436
<212> DNA
<213> Glycine max
<400> 355

agatgaacaa ccaaataaaaa catgacagtg aagaataaaag gagaaataat catttccatg 60
tggataaaag tgagaacaac ttgatTTTGT aattagccta aggtcttaac ttccaataat 120

taagccacct atattctatt ctgaatgact actactcacc aattatctgt acggcccccc 180
ctcccatccg ttcacggata gcacacttgc gtgattcggt gttatatattt actcagcgg 240
cctccgcgca cccctcacgt atgcataatct gatacccctc ggatcgctt atacaccctc 300
cgtccttgcc tctccgtctg actctttcta ccgactatgt atcgtgatga ctcgtattcc 360
gcccgttct cgctcacgag ccactctgct atcttgagt ctcttatctt acttagctct 420
acgtctttat agctct 436

<210> 356
<211> 350
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 356

cttctagttt cactgatact tgtttactat tactggacaa catatctagt tctatgtctn 60
tcannattgn ttattaaaaa gctttcattt gtganaaagt tattatctt gattaatgca 120
ctattcaacc cttcattcta gtgtgatnnt tggtatttca tcatttattt taaaatcgag 180
acatttgate attctaattt aaaaattctg caatcttggt ctccctattt cataatacaa 240
acatntggtc tccatatnta agagaatctg caattctggc ccctatattn tagaaaatcc 300
tcaatcttgg cttaatctt aattgtgtct acattttat ttcttacttt 350

<210> 357
<211> 505
<212> DNA
<213> Glycine max

<400> 357

cttgaacccc atctagtacc cgggatcctc taagccacct gaggcatgca agttgaaca 60
ttctctcatt tgattaagat attattactc tacattttaa gactgagact tatgagaaaa 120
aacacaaaaag ctggggct agactattag cctaggacaa gataaggctt gaagggggcc 180
aagttttatt tgctcaatcg aaaatgcgaa ctaacaccaa tccaatccgg ttatacttat 240
tatgtcaatg aaatcactat taaatcatct aaagtcaatg agatatcgta tgaattgttg 300
ctattaacta acacatacac caaagactag aacaacgaat tgattnacca tcgaatatga 360

agagtgagga gcaccaacaa cattggcagt gtggcataat tttctgcaga cgcatcc 420
accattcgtg tcttcgttt gatggtacat tgctgatgac gccccagag aaaagaaagg 480
attggattga atgaccatca ttgct 505

<210> 358
<211> 460
<212> DNA
<213> Glycine max

<400> 358

caacacccat tacaactctt aatgcattgt ttcttggtgc atttaagcta tcaccccaa 60
caaatttgct taaaccatat gaattcaaac tcgtggtag ttaaccctct tttggaccat 120
caacgttgac ctcaaccata tgtctcttg ttcttacatt agtgagtgaa aattgatttg 180
ctgaattttc attaatgaca ccaatgatgc atctgagttg ggttattgtat tttgtatcaa 240
ctctggtagaa tactgattga tgtaagcttt tgagttaatt aaggtctaaag gtcaatcttc 300
taatgatttt gagctctcca acaataccaa acaaccgtga aaagtgggtg ggtacctata 360
aaagatagag gcaaccctat gaggggggag gtgggtgatc cctttcttt attttctggc 420
gtttccttgc gcttctgttt ctcttactct cgtgttctcg 460

<210> 359
<211> 254
<212> DNA
<213> Glycine max

<400> 359

tagcataata tacaaaccta ggaaacacag attcagttatg ggatacatat atgatatgac 60
atgacaagaa acagacaata tggcacattt tagaagttat acagatatga tatgtatctt 120
aattctaaaca tggctacatg acatgaccac tggttcaag tgtatgtact tcattatcaa 180
gaattatgag aggaagggtg tttatcacag atatgtggca ctatcagata tacaaaagta 240
atgagctatc aatc 254

<210> 360
<211> 114
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 360

gccttgctc tctctactct ctagatgtg gtcatacatn tatcatntaa cgataataat 60
atagagaat ctaactctgg gttctgatat ctgaatcgaa tgctgacatt actc 114

<210> 361
<211> 456
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 361

caagttttn tgataatgac aaagatgatg acaaaaagcc caaagaatga tttcaagaat 60
gagtcaacaa gttcaagatc aagtttatt tcaagttca tgagaagaaa tcaagaagat 120
tcaagaatca agagaagatt gatttcaaga atcaagagaa gattgatttc aagattcaag 180
agaagatgaa ttcaagattc aagagaagaa atcaagaaga ctccacaagg gaagtattga 240
aaagattttt ccaaaaacaa catagcacag ttttgggggg caaaagagtt tttctcanaa 300
natttctaag ttaccagagt ttttactctc tagtaatcga ttaccagttt cctgtaatcg 360
attaccagtg gcaaagttt gatttcaaag tntttactg gaattgcaac gttccaattt 420
attncaaaat ggttatatac aacacaagat attgn 456

<210> 362
<211> 442
<212> DNA
<213> Glycine max

<400> 362

taattgttta attcttactt cttaaatgtt cgttatatac ttgttatagg aaccttataa 60
ttcttaagtat atatagttgt agtatggtgc tctgccttaa ttgcataatgt agtatggtt 120
tttggattt cttgttcata gtgatgctaa tactctatag ctggatgact cattcaatgt 180
tatatttcat aaggaataact ctttgatcg taccttctaa ttcttagtgca acctatctt 240
ttttgtgtt cgtgcttaag tcaaataat ctgattcact tggaaagctg agtataatta 300
attctgtgtg ctatgagact acatcacaca atgggactac ttgatgctc tatcacaatc 360
aagtgattgc tcatgtctta tacgatccac cttgcggta tgcattgttc tcgacttcac 420

gcaatctgct gttatctcaa cg

442

<210> 363
<211> 358
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 363

gtatgtgtat gcagaagatg ataggatgaa gctaacttac aataattcca tgtgctcctc 60
tatagtaact gcttgntata gtcctgaatc gtcctgtcc agctgtatcc cactaaacaa 120
aacacaacan atgaatccaa gcttggcatg catgtggcaa taatcatgaa tatgatcaag ,180
tgaaatgatg atntcgctnt gattctcatt acccaacacc gacagaggtg aatgcagaaa 240
ttatggtaaa ttacaccana caaactanaa acacaataat atagaatcg ggcatattan 300
atgaagttagt cggtaaaga cattgattcg tgtcagtcaa gtgtgatact cacaatct 358

<210> 364
<211> 529
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 364

tgcttgaatc atcgaancct ggatcctctg agtcacctgc ggcacgcaag ctccctccta 60
ttgttgatag agtgtgacat ctatgatcct catagtctcg tgctgtgctc agtgtacgg 120
tagattttgc gcttaagaaa caactgcagc atttgcagct gatatttcta tcttctctct 180
gcattttct tacttgaagg acattaacaa aagccagttt tttttgtat ttttcacaca 240
tttttgtttt ggtatttgac attttgcg cattaattaa ttatattata ttaacataca 300
attagagcat gagagatgga atctattaac agttcaaaa ttttagttt agcctgacag 360
cacaaaaaaag acagaaaattg caaaatgaaa acagtgctaa tattttcttt tctttttgc 420
acgagacata tagttgaatt ggcagtatat ttgcagagaa tatctaaata cagctaaact 480
atctcataat cttatctata atctcgata tagacagtat agataggct 529

<210> 365
<211> 376
<212> DNA

<213> Glycine max
<223> unsure at all n locations
<400> 365

accgtgacca tgtaccttca ccgctaccgt gtacttgatg gtgaccgcac ctctatgagg 60
gggtgaacca ctcggaagag gactgtggaa tacgccacgc tcggtgttgc tactgctntt 120
gtccctccac cctatcatca ccacaatggg tactttggtg ctgccatgcc catggngact 180
tacgttaggg aagcgccacc aaatacagn c tncttccatc accaccacca ccaccaccac 240
caccatgctc gtggaatctc caatgctcat gaaccaaatg ctgcgtccat ataanantat 300
ataattatga ctaggattca gaacaagact nnngatgatga tatgctaact ctcagtaatt 360
ggtgctagag tactac 376

<210> 366
<211> 435
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 366

taactgtctt tgggcttggc ggccatgctc aacaaagtac ttgcacacc tactgtacgt 60
tgatttcacc aatgctgata tggaatgct gctataatct ctatcaaatc ttatcgatcc 120
atgtagatag atattattcc tggtaacca acgctaaaaa ttactgtcta attaattcctt 180
gctccttcta ctattgctag ttccaccata ctttccgt aacttaatcg atgtatatgt 240
ttcgcgtgt ctagctgan tcacagccta agtcacgtat agacaatcta agatgcattg 300
atgatagccc tccacaagat aacgatgact atgcttatgt ttacctactt atcttgaatc 360
taatcgaatg gtgaatgact actatacatt ctatagttac ctcaaattaa cgccaagctc 420
gtctgcagac taacc 435

<210> 367
<211> 468
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 367

ttgagaatca atttgttggc tcagtgtgtg attttcgatc aagccattct tttgttcttt 60

atgttaaagc ttgcattctc aacgttntga aaggaagatt ctgttttat ttcttttct 120
ttcccttac gatcttatgc ttctttgac tctgcattgt nttaactntn ntcttttac 180
tatatactga tgactgtact ttcaatagta ttattaaagg gtaatttctt ccaattaccc 240
ttattaatgg ttaccttcca tctcaggatc aggaantaa ttaataaacat gtttcattcc 300
cacttaatta gcanagnntt ctgaattaac anaaggtna agggactatg tttcttttg 360
tctctcttta cacataagag agtacccctgg tcaataactca actcacgtaa taaagttctt 420
attgaataac ctatgacacc ctataatagc tttatTTAGA tatctaca 468

<210> 368
<211> 351
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 368

tgtccctctc aagagattnt ngattgattt ctatggta agcaccatc ctcttctcct 60
atggccctt gagtttattt tcttctccca ccaagtagac atganatggn gtttcacttc 120
anattatgtat tggttaggtga aaatataatt ganatgagcc tgagtacac caattcatta 180
naatgaaagg gaattgctat ttgcactcct cctttataat aatacaatcc ctatTTTTT 240
atatTTTCC aaaatatccc taanaataca ttcccaatgt tcactccttg caatnntctt 300
tcgtcanatc cctactgtga gtgcgagcaa agagcaacaa tacaccatca a 351

<210> 369
<211> 334
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 369

tagaattcaa ggccctgtaa tccacacaca tgcgtcaagt tccatccttc ttntcacga 60
gtaacactgc ggaggagaat gggcttcgac tgggttgcatt tacactggaa tngagtaatt 120
ccataactgt cggtcaattt ccgacttctg gaagtatggat tatttgtang gtctaacata 180
tngttngct tcaaaacaaa ttatctgctg attaaccaga aatgagaaac gagcaactca 240
ttatataatag gaaataacg ttctgaagat gattcaatta tgaaaaaagg acagattaac 300

cgcatactat tttacctctc tcttcttctt tgtg

334

<210> 370
<211> 215
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 370

ataatcatgc atatgagcat ntcatcttca aaagccaaag actaatgagc tgttatcatc 60
attagaagtt aatattgcat gatcataaca agagagatag tgaaatacca gctttaccat 120
atgctgcac agcaggttgg actttggagc caacagcagc agcaccaatg agcacagtta 180
tagctaaacg gcgagagatt gngctgacat caata 215

<210> 371
<211> 416
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 371

atgacgttat gtttgatcc atttntttta attcatatta agggtgtaaa ataaagtana 60
aagtgtatga aatttttatac tttnnntttt anaatttctt ttaagttaan actggttcga 120
caattttttt ttataaaagtt tagaattatt taaaactaac tgcacactc aacannaaat 180
atattgataa aataaacaac actaagnataa atttttnga aaaattatgt aaaattaact 240
agtaatttag ttgtgcaatg cacaagattg taattaanag gtgaaaaatt gaacaggngt 300
tctaaacaat cgagcactat nttcttaatt ntatcatccn gatacaccac atctaaataa 360
aaggctaaaa atatactata tatatcttag atntcttca tctccatgac atgact 416

<210> 372
<211> 441
<212> DNA
<213> Glycine max

<400> 372

tggagaatag aaagtcgcca caaacattag tatggaaaga taaacatgtt attagggtt 60
tttgtgcaaa tacaaggaaa aatgctatta ccatcttgct cctcttagcc tcttcatcac 120

tgtcattacc atcatcacca acagcttcc tgaaaaagt catagcattc ttAAATgtta 180
agtacatcaa taacaatatg catatcagaa aagtgaaaca acaaaacaaa ataaagatat 240
tggacttcaa accttcttt tgcaatggcc cgatcatcat gaccagcctc agcatggcca 300
tggccatggc tgttagtcagg aaccctgcta acaacgtccc tcagaaagca aagacattat 360
agcttgacac aatgtttct gtcatttcat tcccacttgt acgcttatacg ttgcagatt 420
atagagctt aaacagatac t 441

<210> 373
<211> 333
<212> DNA
<213> Glycine max

<400> 373

tattctatat ttgtgatggt tgcttgcag tgtgtgggc tgaatacatg atagtcgtgt 60
tattgattac aataattatt agtctgggt taattttctt tatgtgcgtt attcacatga 120
gatgttcgct atgaatgacc acatgttgtg gaacaatgtt aacaatgtta gtttatattgt 180
tccgctcgag cataagaggg gtgcaactag acctcaaaac tactgggaga ggaaccttat 240
gagaatctca atcccactat attgaggaaa caacaacaag caagtatcaa catagaagat 300
gtgggagaat taaccacaat gagagaaaat gtg 333

<210> 374
<211> 592
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 374

tacacgtcca ctaccataca cgatgcgtca atctcanatc tgtctaacgt attaacctac 60
annncnnnacc caagacaggt tgattgatgc antctcgatcc cccngatcct atacagtna 120
cctgcgtgca cgcaagctta tgcatagcat agagaggcac ttgggtgtact tgcaactcgc 180
ccttagagctc gcccaggcaa gctgttgctt cactctgaag taacttggtcacataggtg 240
agctgggtac ttttagcccta agccatttgg gggtgcaggt gagtttagagg ctggctgtg 300
cgagccaggg cctagaaaat tggcttaaat gacccttttgc ccccccccccc ttgagtagct 360

tccgcacatctt tgacccaaaac atcgaatgat ctttcgtctt gcgcggtaac tggtgtgaa 420
caactcaatt cagctatcga gaatcacata tccatgaatg atagtcctcg cacgaactta 480
ggcctgacag tgcccccttt acttatttct atcggataa aacgaagtca tattaggcac 540
tattctattt agtgcgctgc tatcactggt caccggcaat ccatggatat cn 592

<210> 375
<211> 347
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 375

gctgctcgcg ttatgcgaga cgtagaccaa catgcttagct attatcgcca agtacctta 60
agagataagg ctggccgcgg cccacgagca taggattgcg ggcgaatatg ctcaagtata 120
cgccgaaataa gaggctagag gaagggtgat cgactctcta caccaagagg caaccatgtg 180
gatggatcag gttgctctta ctttgaacgg gagtcaagaa cttccncgat ttttagccaa 240
ggccaaggcg atggcagaca cctactccgc ccncgaagag attcatggc ttctccgcta 300
ttgtcngcat atgatacact taatggccca catannatag aaatcgt 347

<210> 376
<211> 461
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 376

tcccatagga aggcttaat tntcttgagc agtttgtaaa acaactttac ctttcgacg 60
aacttcaaga ggaacctgga cagggatgct agcctaccat ttagtatttgc gacttcttgg 120
tcattggct ggctgcacat ctccaatatg gcattgcatt cattggggtt ggcttcaatc 180
cctcggtgag tgatcatgaa gccgaggaac ttgcctccgc ctgccccgaa agtacatttt 240
tgaagggtga ggcgcattgtc atactcgccg agctcctcat agacttcttc taggtctgac 300
acatgttggg ctatgctttg agacttgaca accatgtcgt ncacatatac cttgacgttc 360
catttgatct gctatatana gacttggtcc atcagtcgtg ggtatgtac gccttgcatt 420
ttaaagccga agggcatgac cctatagcaa anactggcat t 461

| | | |
|--|---------------------------|--|
| <210> | 377 | |
| <211> | 295 | |
| <212> | DNA | |
| <213> | Glycine max | |
| <223> | unsure at all n locations | |
| <400> | 377 | |
| cttgcgtcct atccttcaac cacttatgat agccaccaac gacanctgtt gctgctcccc 60 | | |
| taagctcctt atctcttctt tccactgcat tccacgcctt gcggattctc tgaagtattt 120 | | |
| ttgcgttgcc ttcatgtaaa ccncatgcaa cgaaaggcgc gatgatctct tccaacggtg 180 | | |
| cccctctcat agggtaacct agttgtctta tggctagtagt gggattataa ttaataacaac 240 | | |
| cccttggtcc catcaagggg tcattcgga atccttcaca tgagcacaac acttc 295 | | |
| | | |
| <210> | 378 | |
| <211> | 434 | |
| <212> | DNA | |
| <213> | Glycine max | |
| <223> | unsure at all n locations | |
| <400> | 378 | |
| atcaatatca tttgtattct tgcataatcct tgcataatcct ttatattactc agtagtataaa 60 | | |
| ataccaagtc agttagaata tggtcatacg gaaattgaat ccagataatc atattatcaa 120 | | |
| gtccagtcag aagatgaaaa taatgtattc tccaatggtt ctatcatgat agataatttt 180 | | |
| aatgtgatgt ttgagtcgtc actatcacct agntgaagga ttggtccgat acatgtacat 240 | | |
| acgttcgtca tcatacgagag tatatcgca cgtcatacgc ttataatact acatgacggc 300 | | |
| atgatccggt tgactaatgg tctgataact ctccttatct atttattatc agtcatcgta 360 | | |
| ctagtcggtt tacagtgttt tataatctga gtgtactgca agtaactcta tccataggat 420 | | |
| gaggcatgta acgc 434 | | |
| | | |
| <210> | 379 | |
| <211> | 461 | |
| <212> | DNA | |
| <213> | Glycine max | |
| <223> | unsure at all n locations | |
| <400> | 379 | |
| ccttgaacgc acataccggt gccaccggag acccccacgt ggtccctcggt gtcttgcacg 60 | | |

ttcaaaccct aatttcaagg ctctacccta cgtntttcac tgtccgaggc tgtctttga 120
attttcgcag attacttggg attcgcgtt cgccagacgg cttaatattg gaaaagccac 180
gaccgttcga cttttcaactc agtagcgctt ccttaatggg tattggcgc taatgcggca 240
tattcggtta cgcatgagaa cgggtattca aaccactatt ggtgcactat tggtagtac 300
tagattgaag cctgagaggt gaaaaagctc acgggctccg cacaactcgt gtgctcaa 360
ctctagtggc ttcttgcgtat acatcactgt tgacttatgc tacactagct acgtctatga 420
gcccttagtc atttccttct tctcgtgctg tgatatgatc g 461

<210> 380
<211> 244
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 380

aatgcgcgca taaacccacc atcccctgtt gccacctcca actgagctca cgtactccca 60
cgtagcccat atnnctcggtt ctctcaacaa cgggtccccca tcaatccttc caagcggtca 120
caacatccaa gaaaaacaac attcatacag cataagctat cacagcccgaa caaaacagag 180
caaaggcaga agactctgct caacacatca accagaatca cagctttct cacttaaaga 240
ccac 244

<210> 381
<211> 431
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 381

tatcgagacg ctganatca tcaacgaaag ctctcgatan attcgaatgg tcataacatt 60
tcactcgat gtccgattcn gngacataat atatcgagac actcgaaagt gaacaacgga 120
agctctcatg atattcaat gctcataaca tttcacacgg atgtccgatt cggggacata 180
actcatctag acgctcgaaa ttgaacaacg gaagctctcg agagattcga atggcataa 240
gaattcacac gaatgttcga ttccggngaca taatataatcg atacgctcga nnatgaacaa 300
ccgaagctct ctagaaattc gaatggtcat aacatttcac tcggatcggt cgattcgnga 360

cataatatat cgagacgctc ganattgaac aacggaagct ctcgacanat tcgaatggca 420
taactttcac a 431

<210> 382
<211> 452
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 382

acaattgatt ttgaattca acgttcaa at acactggtaa tcaattacac catttgaaa 60
tcaattggaa cgttgcaa at ttagttgaaa actttgaaa tcaaactttg cccctggtaa 120
tcgattacag gaaaacttagta atcgaatacc agagagtaaa aactctggta acttataatt 180
ttttgagaaa aactctttg aaaaacaaaa ttgtgctatg tttgttttt gaaaaatctc 240
ttcaataactt cccttgtgaa gtcttcttga tttcttctct cgaatcttga attcatcttc 300
tcttgaatct tgaaatcaa cttcttttga ttcttggaa ac ttttgattt cttctcatga 360
aacttggaa at taaacttgat cttgaacttg gtgactcaat cttgaaatca ttctttggc 420
tttnttcat cattnttggat atcatcacaa ct 452

<210> 383
<211> 270
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 383

ggagatgcag cggaagacan aggagaagag gtgagaggag gcgcattccac aaggaaataa 60
gccatggaag aaggaacttc ggcaccaaga atgtgccttgc gataagaagc ttggagagga 120
tgcttcaatg gagganaaga aagagagaga gaaagagaga ggggggagca tganaatgaa 180
ggaagaaaag agggagagaa gttgaacttt gaagtgtgtc tcacaagact ctcattcatc 240
anagttacca caagtggta acataacttct 270

<210> 384
<211> 173
<212> DNA
<213> Glycine max

<400> 384

tatattgaga cacacaattt cgtgctcctt ctcttcctct ccctccactc atgttctcct 60
tactttaagc tcttatccat gagcttctat ggtggtgagc ttcttcttga ctcatcttct 120
gctagaaggg catctccatc atctttcttc ttttattca ctgccttaaa cta 173

<210> 385

<211> 375

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 385

agatttgttc tgtgaagatc cacagagacc anagctngaa gaggaagccg tcctgagagc 60
tngagaatga tttgtgagtg aatgtgaggt cctagaggtg gaggagacat ccncactact 120
tntgttttg aaatctttca tcttttttt ctctttgtt tanaggaagc ttcccagtta 180
tggaaagcta aatccctctgt tggatcttcc ttgtaggtac ttgatgtaaa tacctgtata 240
tntatntaat gatngtntgt gtgttcaactg tgctatcaga acttcattct accatgctnt 300
ngccttgatc acgttagatgc catgtgtttt aggatcattc aacagtggaa agtggtctga 360
ttcttagaac ttcat 375

<210> 386

<211> 453

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 386

ttagctgttg atctgctttt tgaaaacgta atctttgaga ccgattgttt acaacaggtc 60
aaacaatcga tacttaagga aaagaatggt catgccctct ttcatggaat agtttatgac 120
tgtcaaagaa cgattcatag gactagctcc ttgtgtttta tcaaaagaca aggaaataga 180
attgcacact ctttagcatc tctatcattt tgctatntg ataaatgctg gattcaggag 240
gtccccata aggtggatca natcattca agtaatgtaa tttctgcctt ttcttttgt 300
actggccagg gtcctataaag tacactcgcc aatctacata gcactccatt acgcacgtca 360
gccctccatg tcagccctat cacaagagca cggacgtcca gtccttatta gtcgagctct 420

caatagacaa gttatctccg accatttaat tag

453

<210> 387
<211> 339
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 387

c ttatgttga gaggttaagtggaaaaacagc a agatataaa caaaccatac actnnttcct 60
ctgttctctc ct ttaagac tcanaattga gctgtctcct atctnnntg t tgctgaatc 120
agagattctc tccaccaccc ttntcaagtc acagaatgaa ggtagtgag ggaaaataac 180
aatganacaa atctaaaaat gaaagaaagg acaagtttgt atgtaatgtg aagagatgag 240
agaggaagag ccatatctca tggcatatat tttcgcaaca taaccccact tatgcttcct 300
tatatggag ttagatcata caacacctca actactctc 339

<210> 388
<211> 458
<212> DNA
<213> Glycine max

<400> 388

t tgcacctt g tgatctt gactccatgtc atcgaattgc atgtccactt gtaactcaag 60
agcatcaacc tt tccaac aaaggtttga agaccatcaa acctatccaa aaccttttga 120
agaagagagg aatcttctcc accatgtaaa tgccttctt catcaatggg ttgagcaccc 180
ttttcaccc aagagccatc atgctctta cgataacca aggatgcaat catagtggca 240
ccgattaaga aggatctctt gatttgaaca taaggttcag aatcaggagg gatgttata 300
tgtttaagga agagagtgac taggtgttga tatggcaatg tagcattaa tcgcaatgcc 360
ttatgcatgc gatatcgac taagtgttgc caatcaattt gtcggcctt atgaaaagcc 420
cacatgacaa taagatctc ttcagagacc tgtgcaag 458

<210> 389
<211> 384
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 389

gtcacaccta ctcattctaa aacttaattc cattccanaa cgaccatata tagggaccaa 60
agtacaacat tccaaatcac catctaaaga aaagttcaac ggtgttctac atatgttcca 120
accaagcaca cacagacaaa catgtcatta acacaaatta taagcaaaca aagataggaa 180
gaccgcgagg gggaatgagc gaggganaat gaaccttaca aacgatgaga gagtgaagct 240
attgtgaggg cgagggcatg caatgtatgc gacgataaca cacacgagct tcgacaacaa 300
caactggacaa cttcgacata gacgctntnt gtaacatccc atttttcgt anaaataaaat 360
atagagcana taaaataata aata 384

<210> 390
<211> 449
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 390

gaagaagaag aagaagagga aaagaggaag aaagcctttt gcatatgcaa aatctgaatt 60
gtgaggatta gggaaagggtt tatgaccctt ataattctcc tttcacacc catgcttca 120
ctcaaaacac ccaccccccta acacacatca agacccatca actccaaac tcattgaaca 180
ttatgaaaaa ccaattaatt aattatgaca tcaccacata aataattatt tacttcaacc 240
acttaattta aatttaattt cacaggataa tttttaaaaa ccaattaatc aaacattatg 300
aaaaacacgg tgttacaatt ctcccaaca agaaaatttt catcctcgaa attttcttgt 360
gaagaanata tcgtangcac tgattaagca cacaactatn tcgctttcta ggggtatgga 420
tcttcctct atgatatnct ggatgggan 449

<210> 391
<211> 179
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 391

gtcgaagaac gggcgaaacc ttgcgaaat tcttcacgga aaacgttacg gaaatgttc 60
ggaagcgctt cggcttagat tttcttcacg gaaacaatct ttccaagcan attcgaaaga 120

gagagaagtgcctaaggggc tgaaccctt ccttcttcac ttccctccct atttatagc 179

<210> 392
<211> 308
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 392

atcttcagtt tcggagttgg tcccctagta atntaattca tgaaatgggt cctnctattn 60
tgtanaatcg tgcaatattg atcacccagg ccacaattgg acgttgaccg ttagcaagtg 120
atgttactg tcacgttctt attagatgt gactgtcaaa gtatcggtt ctgtanagct 180
atcttatatg ttgaaagcat gatgttctac aaacatgaca ccaaccagtt aaatcacttt 240
cacatgatan naaagtgata ntttcgtn tcttaattat gcatggataa ntcttatcat 300
tcactact 308

<210> 393
<211> 419
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 393

taatgttgtt ccgagttatg ttggggAACG gttacgaacc cggactgggt ttaggcactc 60
actcacgcaa cttaacttgt ntgacatagg ctaaccgtct tgctggtaa attggatgac 120
tgtctacgac tacatcttcc tttcctttc gctcaatatg gcaacgacta gttattgtat 180
tcgtaacgct cggcaacctc ctacccgaaa gttctctcat tgctgttact actctgtccc 240
tctcattcca cttaatcttg cgtaagtaac tgcctgatct cacttcatta tccggcacct 300
acactacatc agtgcggat agcgattcta tagacgggtt tccttggtgc ggtattcgta 360
tgtccccctc tttgttttc actatcaacc ccactattct ctcttcata ctgtttct 419

<210> 394
<211> 283
<212> DNA
<213> Glycine max

<223> unsure at all n locations

<400> 394

tctctctctc tctctgtact atatatatat atatatatat atatatatat atgtgcgtgt 60
gtgtgtgtgt gtgtgtgtgt gttataaaaa agctaagtgc tgagtgtgat aattntctcc 120
actcatctca aattaagttg gtggtatctc aaatccttaa gcaatgttagt cctanattnt 180
caacaggctt aatatgagag anattcctac aaacagaagt atattgtcaa taattntatt 240
acacataana ttagacagat acatactagt ggtggtccac acg 283

<210> 395

<211> 116

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 395

accgcgatct tagcaccgag ctcagctgtg gacaaacaat agttctttt tacttaatg 60
tttccatatn atatatggta ggtcagaatg tgctttgtta ttaccttacc atatat 116

<210> 396

<211> 478

<212> DNA

<213> Glycine max

<400> 396

caatggctta gtgaggatgg agaggtgcaa gtaggaagc atgtagagtt ggatatttcc 60
attggaaagt acaatgataa ggtgcttgt gatgttggtc ctatgggcc cagccactta 120
ctcttgaaa gaccatggca atttgcataag agagctaattc atgatggttt caccaacaag 180
atctctttca catatcaagg caaaaagata gtgctcagac cattgagtcc acaagaagt 240
tgtgaggatc aaagaaaaat gagagagaaa attttcaag agaagagaga aaaataataa 300
gagagccata cacttgagag ttcaaaaagt caggactaaa ttagggaaac acatgacagg 360
aaacggatga ctgtatcgct tgttagtgagg gacacttctt acctctacta cgatatttgc 420
ataggatcat gatactactg gtcattctgc tccccagttat ttcattttt taaattcg 478

<210> 397

<211> 436

<212> DNA

<213> Glycine max

<223> unsure at all n locations
 <400> 397

gtgctcctta aacctccatt aattnttgc ttacattct cttccattgt ttttcttca 60
 tttttctcc atgtatctcc tcaaattgtct tggctaaat tctgttaaca tgcttcctta 120
 gattttcac ctattaaact tgctatagaa gctaaatttt attttctatg gctcaaattt 180
 cttgctcttg atcttgaacc atgaattgtg ttgagttac gttccttga gtttgc 240
 gatattttt gcccgtgaaa actaaaccat aaaattctta caataatatt acagtagaaag 300
 aaaacctcaa aaatcttagag tgacttgc acctattgtta gttntgtcat agaagtcatg 360
 tctagtcatg aaacttgcataaattt cttatgtgn gctgaatcnt attctctctg 420
 ttcttcgct aactcn 436

<210> 398
 <211> 460
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 398

ctccttttc aaaccatgct atgtgctcgac gactagtcnc tctttccct tcgcagcttg 60
 agttcactat tgctacccca cagagctccg cganatttat tccggccata atcttccttg 120
 cgagccctct tggctctttg ttcaagggtt cttgcgttag ttgcattctc ttcccgtaat 180
 ccggAACACT cttccgaat gtgttagcg gccaacttga acttctccctt ggcaagttc 240
 gcctttctta actcactntt gagagcttgg acttcttcgt cctttccgg tgcttcaaaa 300
 ctctcttcgc tggacttn taacttggtg agccaatcta agcctcgtat atgaactntc 360
 aaccattcat ggtacccacc aatgatgcca ttacgaatgc ccctaagttc ttgatcttc 420
 tttaacnngg gttccatgcc tttatggatc tttgatagtc 460

<210> 399
 <211> 446
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 399

ctccncctt gtcaagcaa ttcttttg atatcatcaa aacctgcatg atttacattc 60
tcccccttt tgatgatgac aaccacctgt aggttaggag caacaagaaa gaaaaaatat 120
ctatttgcata atagttact ccccctgg tttcaatgt ttgcttatata gagacaattg 180
aagatttcat attttcata tataaaaagt tgtctcataaa aaaatagata tttttctta 240
ctattttatc ttttatctt tctctcccc tttgtcaaca tcaaaaacaa atcatgaata 300
gagaggagaa aaaaaatgtt accactgtt gtaatgtatg agaatcaagt gataccaaa 360
ggcattaaac caatcattca atattgatca agcaaaaaca agtatagtaa cacatcaatc 420
anaaaacacaa tcaaaagcaa tcaact 446

<210> 400
<211> 469
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 400

cattgacaac atgattgacc cttggcagc cgaagttaa tgtgccaagc tatgccttt 60
ggggttattg ttgctttga aatttaacc aaaaatgact aaagttaggt taaacccaaa 120
atggctaaaa tggctaaagt aggttaaac caaaaatggc taaagttaggt taaacccaaa 180
aatggaaaaat tttgctttt taaaactgg taaaccctat cataatcccg tagatggatg 240
tgctaaccctt cttggatgt gtaatcagag tgaaccttgc acaaagtcca ctctcacaaa 300
gttaaattac atagtcactc aatgcacaat gcaattctt gatagataga aattcagctt 360
agacaatttt catatctcta tatcaaacaa aacacatata cttgttata tatctatgt 420
tagcacgaaa ttcaaggctt agaattatac aatntataaa taaattct 469

<210> 401
<211> 451
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 401

ctaaagaagc tagagcttag ctacacacat ccnctataat agctaagctc acccccaagc 60
caaaatacat gaaaatataa aaaaaagtcc ctaatacaaa gactactcaa aatgccctga 120

aataacaaggc taaaacccta tactactaga atgacccaaa tacaaggcct aaaaagaagga 180
aaaatctatt ctaatattta caaagaagag aggatccaac ctgggtccat gggctcagaa 240
atctaccctg ggattcatga gaaccncaag gccttcttta gcagctctag cccaatcctc 300
tttagagtctt ctatccaata ccccttgtgg ggtaggatng cttcattccc ttcaacttgg 360
aaaggatntg acctgaaaat ccgaaggctc tcataatttgc ggctccctcc ctcgacacacct 420
cgaaaaaaaga ataaaacata tgtatttagtg g 451

<210> 402
<211> 380
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 402

gctgaatata tagctgcagg aagttgatgt gctcanagtc tttggatgaa gcaataactn 60
taacactctg gagtanacct tgatcacatt cctctaaaat gtgacagcac aagtgcgatc 120
aatctaakan aaaaaatctt gtcatgcatt cttagaactaa acacatagaa ataaggcatc 180
attntctcaa agatcatatg ttaaaaagttt attgggtcat tgagttcata gatagtggc 240
atcaactaac agacgttncc actanaccac ttgcttagaga tagattctnt ttcgttagaa 300
atgaactaga catatttagat gcatctagta tagaatgaca ttcttattgc atagtgtgtg 360
atgcacattc ttactcatat 380

<210> 403
<211> 114
<212> DNA
<213> Glycine max
<400> 403

ctaaaaactt agtagataa aggttataga tttaggcgac agtggcagcg gcaattatac 60
aacgcctgcc tacataaaac tactgattct acatagatat ccaggacgat cacg 114

<210> 404
<211> 393
<212> DNA
<213> Glycine max
<223> unsure at all n locations

<400> 404

tcagctgaat cacttccctc ttcanacact agaatagata ttatatnngg tctttataac 60
acaagtctata cctaattcng attggctctga ctaaacaatg tanaagtatt tatacttagtg 120
atttatgaaa aataaattca ttntanatnt gtgttaatt ntaatttatt gctggtgtaa 180
ataattttac cccatcaact aattaanaat cttaatgtaa ttataagaat ttaatttatt 240
agcacccaca atataatata taggaaccag tacaaaaaat tcttttaac attagttgn 300
tttaacattn tataatatgtt ttgtatgtt tgctaataag ttgatataga aacactacaa 360
gaaaaacact taaacatgtt tgatataattt ttt 393

<210> 405

<211> 241

<212> DNA

<213> Glycine max

<400> 405

ttatcctgct ttgtatgata tgaaggctcg ggaagatgga gagataagaa agagggagaa 60
tcatgttgc actgcccct acatggccaa attccacaac taacaatgca acacttagct 120
agatagtcat ttcataccca ccacctacta gtaagacact tatcatcaca aggccacctt 180
aatcagcaca aagtcacctg ccgacatcta tataaacacc ctcttacact accaaacact 240
a 241

<210> 406

<211> 246

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 406

atcatgataa caaccaatat gaattccact canaggagtt gggcatgtaa aagccaaaac 60
ttcttcaagc tntagcctta agttgttcac catgttgctc ccctatctct aacaacccat 120
gcatgttagtc caagttcaaa ggattatagt atgttgatag tgggcgcata aaccatatga 180
taagggactc aagtctgtta aactctttag acaaggctgt tagaacccaaa gtcaagaatg 240
gaaatg 246

<210> 407
<211> 454
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 407

gatatccact ccacaagggtt tgaagttagag gagagcttca accctataac gcaacgtggc 60
ggacaaaagt gggcagtaaa cttgaatgtat cgtcattgtc aatgcggaag gtattctgcg 120
cttcactatc catgttcaca tattattgca gcttgtggtt acgtgagcat gaactactac 180
caatatatacg atgttgttta tacaaacgag cacatcttan aagcttactc cgacacaatgg 240
tggcctcttg ggaatgaagc ggcttattcct ccttctgtatg acgcatggac acttatccct 300
gacccaacca caattcgtgc gaaaggctgg cccaaatcaa caaggataag aaatgagatg 360
gattgnngtcg aaccatctga gcacccgaaca naatgcagta gatgtggagc cgaaggccat 420
aacaggcgtc gctgtccaat gcaatctgag cgtg 454

<210> 408
<211> 304
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 408

ttagctatat gnggttactt atatttgtca tggatgatn natttagctt tatataattgc 60
taattcagct tataaggtagt aatgaagcta ctntagctcc tatatatnna accatngatn 120
ntaatagcaa tatggcactt ntgagcaaga attactctt ataagttntc atatcaagag 180
tcanatgcta ttgaaatgga taaatgcaca atataattgg tgtgtatcaa ccctaacaca 240
acaacactac cacaaaaaca cacacnctat gatccacaat tngaaacgaa agggaaaagt 300
catg 304

<210> 409
<211> 457
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 409

aaaggatcga agtgggtctg aaaagaggca aatttaatca tcctgcttgg gcgaatgaga 60
aaactggggc aattgaagaa ggtgaggatg agggagaaaac ccatgctgtg actgccattc 120
ctatacgacc aagttccca ccaaaccaac aatgtcatta ctcagccat gacaaaccct 180
ctccttaccc accacccagt tatccacaaa gccatccct aaatcaacca caaaggctgt 240
ctaccacact tccaataacg aataacactt ttagcacaga ccanaacacc aaccaagaaa 300
tgaattntgc agcgaanaag cctgttagtt caccccanat tccgggtgtca tatgctaaac 360
ttgctccat atctacttga tactgcaatg gtagccataa cccctactan gtttcctcaa 420
cctccatTTT tccgagggtta cgactcgaac acaatgg 457

<210> 410
<211> 254
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 410

tagatagatg acatcnagcg ataactgagc catgtaatg ctcatncgtg tcaggatggc 60
gtacactagc taacacttcg gcagggggag gtcggaatta tgatcgctgg gctggatgtt 120
gctgagcagc anaatcatcc agatctgagt cagggtggtc atggtggtgc gcatgatccg 180
cacncgtctt ccagcagcag tccaggcana atcctgcccc ggtatacata gcaactggc 240
gatggccctct catc 254

<210> 411
<211> 455
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 411

aacctatatt taaaataact taatgccatt aacctaggaa attaaaaacaa acttaatggc 60
tgagtgtaac tgatattgtg gcaaccaaaa gtcacccca acagccaaaca agtcagccac 120
catttggtct cccaaaaggc tgatgcctat gttgccaatt gggcccttat tacaacttga 180
actaaagccc ttttagttga ttaacccaaa acatatttt ggtcagccaa cttaacaagg 240
attggccat tat tagaca aactaaacac tctaaaactg aaataaagtg gtgtcattta 300

gtcctcctcc atttgggcca tgataacaact cacaaccttg gactttctc cttganactt 360
gngcttgtat tcaaatacgta tggacagcac tttgtgaaga gcttcottgg cttdcccttgc 420
tctaccctt gtcataangtc ctccaaaagt ctan 455

<210> 412
<211> 352
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 412

tgaggtaatt ctctctctct ctctcaatag ctgaggaaaa gtagttccat gaagaagatc 60
caagccgagg cgottccgta acgtttccgt gagtaattac gcgaagattc tcgaccgttc 120
ttcaagattc atcgctcggtt ctgcgtntc ttccgtcttc aacggtaag tacctcagac 180
caagctnttc aattcattct atgtactcgt ggtggccaca ttntgttcat gtattttatt 240
ctcgtttcat ttacttttat acccgctttt acgtgcttaa gccattattaa agcatttctc 300
gctaatactaa aataaataaaa ttccaccgat cgttgaatgg tatcatcgta at 352

<210> 413
<211> 448
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 413

gtcaaggcag ctgaaggAAC tagttccgct ccggagtatg acagtacccg cttaggagt 60
gttgtacacc agcagcgctt cgaggccatc aagggatggt cgtttctccg ggagcgacgc 120
gtccagctca gggacgatga gtatactgat ttccaggagg aaatagggcg ccggcggtgg 180
gcaccactgg ttactcctat ggccaagttt gatccagaaa tagtccttga gttttatgcc 240
aatgcttggc caacagagga gggcgtgcgt gacatgagat cctgngtaag gggtcagtgg 300
atcccgtttg atgccgacgc tatcgccaa ctcctatgat atccgttggc gttggaagag 360
ggccaggaat gtgagttatgg ccagaggagg aaccggcttg atgggttcga tgaggaggcc 420
atcgccccagc tgctatgtat acttgggg 448

<210> 414

<211> 278
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 414

 actggtaatc gattaccana acattgtaat cgattacagc tnnnttgaaa taatcgAAC 60

 gttgtaaatt cnagttgaaa aactttcan aacaatattg ctactggtaa tgtcataACCC 120

 taatttcgtc cggggacctt tgcttgatga catgcgacct ttctttggTC cttgtgaggT 180

 gcttggcatg catcattang caatnTgtGA gattccAGGA catGCCGACA. aaccaacaaa 240

 atattgatgc acaaATCCGT aagtttccgt gacacACC 278

 <210> 415
 <211> 457
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 415

 cgggcattc caagtgttgg agaagatcaa cgacaatGCC tacaagATTG acttgcCTAG 60

 ttagtataat gtaagtGCCA ctttcaatgt gtctgatcta tctcttttG atgcagatGG 120

 aggagCCTTg gatttggAGGA caaatCCTTt tcaagaAGGA gggagtgtATG aggacataAC 180

 caagggcaag gaccatgaag cacttgaagg tcccatgacc agaggcAGAC ttaaacaAGC 240

 ccaacacGTC atagagacAA ggctggTCat ttgtatAGCT gccattgtATG atgattGAAG 300

 gcccAAGTGG agaaAGATGA aggcccAGAG gcagaggcAC taccaAGACT actaATTGTT 360

 gttgaaggCC CATACTAact tgaaggCCCA agttaaATAA gttnttagTT ataatttATT 420

 tntattggaa ttctggCCCA tactgtntAG aacGCCN 457

 <210> 416
 <211> 511
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 416

 ngtgtgctcc tgctttacg gacctatgaa tctcagctnt cacaagAGTC ttcacAAATA 60

 actatcatga agcaganaAC taacAAAact acccatcata tctccccAAA ccccatACCC 120

acaaatgg agttcaatg gagaatgaag aagaagagaa tggcaacgtg agggagagag 180
agagctgtct gaaataatgt ggggctgagt gaagagagag agagttgc ttgatttt 240
aaaaggctt ttccctcattt ctttattt tattataaac tatgccacat gtctccattt 300
gagtggagca aaaaggcccc actttccctt ttgactgtga cccatactca gccacaaaaag 360
tgagggaaat ctgacctttg aaatgctaaa atcctgcctt gttggcgtg ccgttctct 420
gttccagtt cctcgcggtt ctctgcgtcc atcggncca gtttcgaaa gtacgcaata 480
tatatatcan aacgctcaga ataaaacccc g 511

<210> 417
<211> 445
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 417

attctattag cacggattnt atcagctcaa caaaagttt tagatgcaga acttgttata 60
gatgctgctc tagatcaaag tggcaaattt gatcaaggag aattgtttag aactaaagct 120
aaactacgga ttgcacaagg aaaattaaag aatgcagtgg agacatatac ttttcttctt 180
gctgttcttc aggttcaaaa taaaagttt cgcacagcaa gtaaggttgt gaaggtatgt 240
gtcaatgata aatgtagaat atatcactta gattctttaa taactcanat tatcaacggg 300
aaaggaatat tagaagcttc gtcacagtc ttanactcta ctttttaca tatgtagaat 360
aatatattac ttttctgatt tatcttcat acagaataan ggaaaccgtg acagaagact 420
ggaaatggaa atatggcttg attat 445

<210> 418
<211> 480
<212> DNA
<213> Glycine max
<400> 418

ctcttatttc ttggtaaagc tctatctaaa taaagttttt attactgcaa gaaatcagat 60
ttatctccca cccctgtcct tctcaatcat tccaaatcccc accccccaaa gtccatgagg 120
gtacccttca ttatgtggtg cttacggctt atacaaatct aataatttgt ggattcaggt 180

agatgtggag ttttgatgg tactattgaa aacatgcact tgcactggaa gtactggaa 240
ctggtcaaga aaattgtgaa ggctaaaacc tttgaacgag taaaaaaaaat tgcactagca 300
cttgaagctg agagtggagg tttctggc tcagttgaca aagtttcaaa agatattct 360
gtaattgttt atcggcgtaa ggattaccaa cgtccttcaa cattgagacc cacaaatctt 420
ttgacataga gaaaggctt agcacgttca atcgagcttc aacgacatga ggtatgtatt 480

<210> 419
<211> 460
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 419

aatggagttg acaagaatat ctccagactt atcaacacat gcacagtggc caaggatgca 60
tgggagatcc tgaaaaccac tcatgaagga acatccaaag taaagatgtc cagattgcaa 120
ctattggcta caaaattcga aaatctgaag atgaaggagg aagagtgtat tcatgacttc 180
cacatgaaca ttcttgaat tgccaatgct tgcactgcct tggagaaaag aatgacagac 240
gaaaagctgg tgagaaagat cctcagatct ttgcctaaga gatttgacat gaaagtca 300
gcaatagagg aggcccaaga catttgcAAC atgagagtgg atgaactcat tggcctt 360
caaacccttg agctangact ctcggatagg gctgaanaga agagcaagaa tctggcggttc 420
gtgtccatg atgaaggaga agaagatgag tatgacctgn 460

<210> 420
<211> 269
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 420

accatacaga accttgcct tccatgcagc aacctggagc aattgagcaa cctgaagctt 60
atgttgcana tatttacaat agacctnctc aacctcaaca gcaaaatcaa ccacagcaga 120
acaatttatga cctctcttagc aacagataca acccttagatg gaggaatcac cctaattctca 180
gatggtccag ccctcagcaa caacaacagg gggtagggaa agtacccct tgaattgtat 240
attcaagaca tttgagaata aacaaacac 269

<210> 421
 <211> 427
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 421

 ctaanaagag gtatatgtga tgtgagagaa aaaaacatac tcatactccg attattaaca 60
 gatgcttct ggtgttgcac acgagttgcg gcaggatgag ccaagcttc aatggacc 120
 atcaaaataa gaacattgaa gcttctaaag agtgacacgt atattataca aatataatag 180
 tttagaaatag atagtatcat attatacgat atatatatca gatgactaac ataagatgat 240
 cactgcttagc tggacggcag caganaattc atgccaggaa acgattaaat tttgacttta 300
 ttaattcttc tagcaccta taatggaaaa aagagttgat agattgcgc ctaaacattt 360
 tatttaaaac aaacagagtt tccaacatcg attgagagtt ttttatatc aaacctgtga 420
 ataatgt 427

 <210> 422
 <211> 365
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 422

 cactcacaca ccgcactctc caatccttag ctcanaccta tggcccttg atgctacttc 60
 actttggaaa aatgccagtt ctgggtgtct caactgctga ngctgcacgt gagggatg 120
 aaacacatga cctcgtnnc tccaacagac cacatcgtaa gatgtttgat atcctttgt 180
 atggttccaa agatgtggca tcttctccat atggcaacta ttggaggcag ataaggagta 240
 tatgtgtctt gcacatcttc agtgccaaan agttcaatc tttgggtgca gtgagagaag 300
 aagaaatctc cataatgatg gagaagatan ngcagtgtgt cttcttgatg ctgtgaatta 360
 tctga 365

 <210> 423
 <211> 448
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 423

ttcanagttg ttttcatatg aacaatgtat acaacaaaag atgtatgacaa aggtatgaa 60
 aaaaagctca aagatcaaag aacaactcaa gtaaatcaa gaacatctca agtgaatcaa 120
 gaacaagtca agagttctag aatcaagaag aattcaagac ttaagaagaa agcctagaat 180
 caagaatcaa gattcaagat tcaagaatca agactcatga ttcaagaatg aagaaaagac 240
 tcaatcaaga taagtattaa aaagttttt ttttaaactt tgaatagcac atgagtttt 300
 gacaaaacct ttaccaaaga gttnttactc tctggtaatc gattaccagt agcaaaataa 360
 gtttgaataa gtttcagac tgaatttaca acgttccat tatntcaaa aggctgtat 420
 cgattacaat gttntggtaa tcgantag 448

<210> 424
 <211> 469
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 424

ccatcaaccc catgccaaaa tacatganaa tatataaaaa aatccctact acaaagacta 60
 ctcaaaatgc cctgaaatac aaggctaaaa ccctatacta cttagatggc caaaatgcaa 120
 ggccaaaaag aaggaaaaaa cctattctaa tatttacaaa gaataatgga tccaaccttg 180
 atccatgggc tcaaaaatct accctaaggt tcatgagaac cctagggcct tcttttagtag 240
 ctctagccca agcctcttgg agtcttctat ccaataccct tgggtttagg attgcatcac 300
 accatacaac attggttttg accatcaatc actatccctt tgtggttgat tcaccttcaa 360
 atcatattta tgtttggaaag agagaaattt ttgttgtct gagcgtaact totcattctt 420
 tgttgatctt tcacactcca cttcacctt cactaatcaa ctcaaacct 469

<210> 425
 <211> 234
 <212> DNA
 <213> Glycine max

<400> 425

tctgtgacac catcagaccc atgccttcat gcagaacctg agcaataggc agccgaagtt 60

atgctgaaat atttacacag acctctcacc ttagggcaaa tcaccatgca aacaatatga 120
ccttcagcac agaacaccct gatgggaat acctacctca aggtcagcct cacacaacag 180
agctgctttt cttcaaattgt gtggccagag acatcattct cacatccaca caca 234

<210> 426
<211> 418
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 426

gctacaacaa ctgcagcaag acccaactatc cacaatgaca gaacaatatt tgtntatacc 60
ttgcacatctt tatgaaagat gttctcttt tgggtttggg ttaggtcaca agattgactc 120
ccaagtaacc ttctgaccat tagaagatca cttcttcat agggtaaat ctcttcaata 180
tggtcatcac cattggcttc accctcactt ccactngagg aaggagaaga tgtagcctnc 240
ttttggctac tatacatgtc ttgaccgctc atgatcatgg ttttcttgc gggcaatga 300
gaagcaatgt ggcctagcaa tgcttgcctt ttcctcctcc ctaagtctag ctctcagaag 360
ggagtagtgc atttgcgtt atattctca cactcatact ccctngcta agctttt 418

<210> 427
<211> 589
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 427

ctaacggctt cctccatatg tcataggcgt taaaagcggtt gtaatatctc gtaacacttc 60
aaaannctta gcactagttt gaatcatgcg tcgctgagac ccntagagtc gacctgcagc 120
atgcaagctc tgcctttaga tccctttgt tggactatac tcataccaag caacattatt 180
gtacaacata ttataaccaa cacttaatcg gcagattcct ctttagcagac taagattcaa 240
ttctgcttca ttcaaggctt aaggcaacaa tacattttcc aatgctaaa tcacctaacc 300
gggcacacaa atggttgc agaccatgag cataaaaaat ttaagcactg aaagaagcat 360
tgaacacact agaaactcaa tcaatttagat attaaaataa ttacatcagg tggcttttag 420
aaatacccaa caagggtgtt tagcccacca ttacagacaa acccctatca ataatgagat 480

aaataaaccg taagatttct tgaaagctgt cctttgctt ctacagagct tttccaaaa 540

ggcacttggg tgcttataat ttgtgcaaag cgtgttaat ctggagtan 589

<210> 428

<211> 192

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 428

caaccaaaga tatgaagatg tgagatgtnt ggttctctgc catcgaataa ttcatatgaa 60

gtnttctnta aaatgggtct tattaaagcc ctatctaaaa tgttagcatgc agtgttaacg 120

gcttcagccc ataagtattc tggaagagga gtatcattca ataaagnctt agcaatctcg 180

tccaaagatc ta 192

<210> 429

<211> 526

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 429

cgttgatcgc tcgtgacgag ccnnnttgagt aacatcgacg catgcaagct tgcatatgaa 60

gaanatgggt gggtgggcat taaatgtgt cattaaatgc atatacttcc tcatgctaag 120

aaactagtct ttgtcgcagc gtattgaaca cttaaggagg aaaccacctc tttgtgtta 180

gaacaagttt atgcaacaaa gttcttttt tgatggcgat tgagaaattt tagagcttga 240

cttcatttat tcctcatagg atgtgacaaa tcctaggaga atatctctgt aaaatagatc 300

tcaaacacaa gagtattaaa tgaagtctt catgtcaact ttaatgtgt atcagatcat 360

gatttcatct tggctaccat tgaaacatca gatcgacact ntgcaaaaca tgatttgata 420

gcacataaga tgcaactnta aaccttcgta tttgtttca tctaacaaca tgtcttaaga 480

cataaatgtc ttttaaccta gcaggaatgg tattcctatg aatacg 526

<210> 430

<211> 250

<212> DNA

<213> Glycine max

| | | |
|---|---------------------------|--|
| <210> | 433 | |
| <211> | 456 | |
| <212> | DNA | |
| <213> | Glycine max | |
| <223> | unsure at all n locations | |
| <400> | 433 | |
| gaaacttcct gcttttattc gttgaccaca gagtggtacc tggagatatg tcgcggnggt 60 | | |
| caggagacct tgnngacgta aggtgggtg ctattgccca aaaccaagct tgaccaatcc 120 | | |
| cgacccaacc cgggcatagt cagtcagtga gagcctgtga tgtacctaata caggcgagct 180 | | |
| cctggcagtc aacagataaaa aggaacaaag accacaaagc aaggaggctt gtggtggtcg 240 | | |
| gccagctgtg aactttgatt gatatgtgag atttggcctc tggtaatcga ttaccaaggg 300 | | |
| tggtaatca attacaaggc ttanaaatga agacagaagg ctaagatggc ctctagtaat 360 | | |
| cgattaccaa ggggtgtaa tcgattacca ggcttgaaaa cgaggtcagg aagccatgan 420 | | |
| ggcttcttgtt aatcgattac caaggggtg taatcg 456 | | |
| <210> | 434 | |
| <211> | 318 | |
| <212> | DNA | |
| <213> | Glycine max | |
| <223> | unsure at all n locations | |
| <400> | 434 | |
| gaataaagag ggggagaagt agaactttga agtgtgtctc ataagactct cattcatcan 60 | | |
| agctacaaca agtgttacac atgcttctat ntatagacta ngtagcttcc ttgagaagct 120 | | |
| ttcttgagac aacttccttg agaagcttct ttgagaaaaac ttccttgaga agcttagagct 180 | | |
| tagctacaca caccctctc ataactaagc tcaccttctt gagaagctt cttttagaa 240 | | |
| ttcgtagata agcttagagct tagctacaca tacctctcta atagctaagc tcacctcctt 300 | | |
| gagatgagaa gctagagc 318 | | |
| <210> | 435 | |
| <211> | 215 | |
| <212> | DNA | |
| <213> | Glycine max | |
| <223> | unsure at all n locations | |

<400> 435

tatgtggact aggtggcgat cgacgatgg tgcaagtcga ctcttacat ccacaaatca 60
cacataaaatc catcatcccc agntggccac cttcaactga gctcacgtac tcccacgtag 120
ccncttatcc tcgntccttc aacaccgggt gtccatcaat ccctgcaagc ttccacaaca 180
tncaaggcaat tcaacattca tacatcatga actat 215

<210> 436

<211> 308

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 436

gactganaat gttattcagt ntgtcagaat ggatgtgaag ctttgattt gttggcagcc 60
caacccatt acanacctag aaagtccttc ggattcattn tgtgtgtta tttctgtatg 120
gtatgagatg atatgcanaa gttggactt ctgttagttg tttataatgg aatgatccta 180
aacacttgag cttgagtgaa acaacgactg tgaggctng gttgatgatt ctcccttga 240
tatctgccat tctcaactaac ttattnagt tgtgactcta atgcataatgt tcctatcatt 300
aaaaagct 308

<210> 437

<211> 456

<212> DNA

<213> Glycine max

<400> 437

actaacgtcg tcttctgcga cctttgtcaa tcgcggccga caagcccggt gacacgtgga 60
gatttacgtc atcttcccgcg ctcacaagat ctgtcatact gacttttag tcacgctgac 120
ggccggaaat atccgagtgg ttatccgtat aaacttttg ctgtctgtaa gacgaaaagc 180
ttgatagcac gcagagacta acgtcgctt ctgtgccatt catcaatcgc ggccgacaag 240
cccggtgaca cgtggagatt tacgttatct tccgcgtca caagatctgt catactgact 300
ttttagtcac gctgacggcc ggaaataccc gagtggttat ccatataaac ttttgctgt 360
ctgtaatacg aaaagcctga tagcacgcag agactaacgt cgtcttctgc gaccttcgtc 420
aatcgccgccc gacaagcccg ttgacacgtg gagatt 456

| | | |
|---|---------------------------|--|
| <210> | 438 | |
| <211> | 254 | |
| <212> | DNA | |
| <213> | Glycine max | |
| <223> | unsure at all n locations | |
| <400> | 438 | |
| actcanaagt cagtatgaca gatcttgtga gcgcggaaga tgacgtanat ctccacgtgt 60 | | |
| caacgggctt gtcggcccg attgacgaat gtcgcagaag atgacgttag tctctgcgtg 120 | | |
| ttatcaagct cttcgtctta cagaatgcan aaagttata cggtataacca cttcggatt 180 | | |
| tccgcccgtc agcgtgactc anaagtcagt atgacagatc tttttagcac cgaagatgac 240 | | |
| gtanatcacc gcgt 254 | | |
| <210> | 439 | |
| <211> | 487 | |
| <212> | DNA | |
| <213> | Glycine max | |
| <223> | unsure at all n locations | |
| <400> | 439 | |
| atgtataata aaataggtaa tgtgatacat agaaggatga caagcaataa aatcgaacat 60 | | |
| aagaaaagt tacaaaaagt acgaataata ttaagataac ctcacataat ggagaaacag 120 | | |
| cattcttgat aacaattcac ttccatcaca agaaanaaga tctgataccg tggactgatc 180 | | |
| aacgcatac tnganaaaaga tatagaatag ttatatctt gattcagtgt atggccaaa 240 | | |
| attgacggta cagaatgtat gaagagagtt tagtctaatt aactaaacag aatataccaa 300 | | |
| tattgtaaac tntgttatgg tgttcagcta gtacggataa ngaaacaata caaaatttga 360 | | |
| tctaaataat atagctctta tgtcaaagca caatangatg attttaaca aatgactgaa 420 | | |
| tcaacacgca tatatttac aatctccaca aagatagaga tcataaaac atctcttata 480 | | |
| tttatat 487 | | |
| <210> | 440 | |
| <211> | 439 | |
| <212> | DNA | |
| <213> | Glycine max | |
| <400> | 440 | |

ttgtcgtgat taccaagtga cagaacaaca ttacctgtat tgagcaactt caataagaac 60
ctttacaggg atgctatcct accctttagg atttggactt cttggccata ggtatcgctg 120
cacatctcca atatggcatt gcatttattg tggggctt caatccctcg gtgagtgacc 180
tgaagccgc gaacgtgcct tcgtctgcac ctaaagtcat tggtgacagt ggcagccat 240
gtatactccc ggacatctca tagacctt tttagccccgc aacatgtagt cctctgcttt 300
gtaattcgac tcctatctag tgcaaataata ccttgacgccc acattcgatc tgatatttat 360
aggcataggc tcaacctacc gtgtgtatgt caccgctgac ttatctagcc ccacgcgcta 420
ccctaaacaa ccctcgccg 439

<210> 441
<211> 419
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 441

caacaagtat tctncatcat gcacacac acantcatat caagagaaaa aggacattaa 60
tcctttccc ttccctanggt cacaatcaac atggcccttc aagtggaaa atccacttta 120
ccaaatcaca caccaccaca tatccaatca ccaaattcatt actagacatt caaagtanna 180
ttttctgaa gggtggacac nctttgacct aaccctanag tgcgacgaat cttaaattat 240
atcattaata aactcatata cataacacac aacattctt accaagtgg cacactcaat 300
tggtcttaan acatatacaa tagcaattct tataattca tattataaaag tctncatcan 360
agtanacaat acattctaca attcacatca ttaatttcat gcatttcata tacattcat 419

<210> 442
<211> 463
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 442

tcaaaggcctt gtatggattt aaataagcta taagagctt gatatggaaa ctaagtttat 60
tcttactcta gatagttact ctagagaaa agtggacact tcactattca gaaaggctta 120
gaaaaaggat ctgctgatta tacatatata tgtgaatgac atcattttt atgtaacctc 180

tgaaaggatg agcaaggagt tttctgagct aatgaaaaga gaatgtaaaa tgagcttgat 240
ggtaagttg aagttcttta tangactcta aatccttcaa caagattatg gaattttcat 300
gcataaagag aattcctcat ggaccttattg ataatgttat taacggatga agtcatacat 360
atggttaccg tgccatcctt ccactatctt tactatctaa tcattggat gctatcttta 420
tatgaccact tactgtctat atcgctcctt tttcttaatc ccc 463

<210> 443
<211> 433
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 443

aaggtagt ttgttagggct ctctctcttt nttnnnnnn ggttttattt ttagagtgct 60
actctcaatt cgtttagtc tctttcggtt ttgaaagaat aattctagtt ttcttcattt 120
tctactgatt aatggaaggc taagtctcca gcgttatttt ctcttgagga ttaagcacaa 180
ctctcttga ggttctatta ttactattaa attctgataa gttttccctc ttcaccaatt 240
actatgtatt tggtgctatt aatccatgca tgcttagtgc ttgattaatt gtctctgtgc 300
ttaatttacg ttcatgctta ctgatcggtt atgattaatt ggtgtatgtg ttggtaatc 360
acataatgaa tgccttatgt taaatttgc tttagtaannt taattanggt tggattaagt 420
ggttgaactg acn 433

<210> 444
<211> 161
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 444

tatgaccttc atgttatggg tatccttgc gacgcatatt angntaagag ctcttgtat 60
aagcaacact ccaccaagag gggaaaatgg tgaatggagc attttggat tgtaacataa 120
gtatatgcgg aacacgaaga aagcaaggta tggatggag a 161

<210> 445
<211> 371

<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 445

ctcagcta at aaggagga ag tggagagg tt ggaagctact ttggccaccg cccagaggga 60
gaggaatgct gaaaaggagg agaggattgn ggtttagcaa aaaacatata acaatgttta 120
tgaagcttat gagtggtgggt tgaatcactg catctaacaa ttacactttg actatgaggt 180
cccaaatgaa tccat tttca acatgaacaa agatgtctat aatggagact taatcttgat 240
tcatgacatt tcggatgagg tggatcaaa tggtggcag ccaactaccc ctcttggta 300
aacccaaatg accaattcg ctgaggagga tgtggacgt gtcccatata cgagccanaa 360
tggcg tctct t 371

<210> 446
<211> 436
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 446

tg ttcgcaca tcg ttcgcgt gtatgatatc cactcgacaa gg tttgaagt agaggagacc 60
ttcaatccta taacgcaacg tggcggacaa aaatggcag ttaactt gaa taaccattat 120
tgtcaatgcg gaagg tattc tgcgcttcac tatccatgtt cacacattat tgcagctt g 180
ggttacgtga gcatgaacta ctaccaat atagatgtt gttacacca tgagcacatc 240
ttaaaagcat actccgcaca gtgg tggcct ct tggaaatg aagcggcaat tc tcc ttct 300
gatgaggcat ggacactaat ccctgaccca actacaattc gtgcgaaagg tcggccaaaa 360
tcaacaagga taaggaatga gatggatngg gttgaaccat ctgaccaccc acaaaaatgt 420
agttagatgtg gagctg 436

<210> 447
<211> 442
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 447

agcgcgggtc tgggagacaa aggtcaagcg ttcgcgatat gcgaggatga tattccgagt 60
actttggatt tggtacgacc atgccctcct gatttccagc tggaaattg gcgagtggag 120
aacgcggccgc gcatttacgc aacgagcata atgtaaacct ttacggttt aaaagctcta 180
tagttgggcc taggccttag agttttcct ttgttaagg ctttgtgtct ttgttttt 240
aatttataat acaaggatct ttcttcatct gttcctggtc tctaccatt ctcattcatt 300
tgcatgttta ctctttntc tgaaacggca gatccgatga cgagcccccc gaaggtacta 360
atacctggga cccgcctatc gacttcgagc gagaaatgaa tcanacggaa gatgaaggaa 420
atqaggatgt qqqacttccc cc 442

| | |
|-------|-------------|
| <210> | 448 |
| <211> | 410 |
| <212> | DNA |
| <213> | Glycine max |

<223> unsure at all n locations
<400> 448

ngagttgagg aagtgttagaa gggtgaaact tcctgctntt attcgttgac cacagagtgg 60
tacctggaga tatgtcgccgg ngtcaggag accttgnnga cgtcaggtgg ggtgctattg 120
ccccaaaacca agcttgacca atcccaccc aaccgggcgt tagtcggtca gtgagaacct 180
gtgatgttacc taaaacaggcg agtcctggc agtcaacaga taaaaggaac aaagaccaca 240
aagcaaggag gcttgtggtg gctggccagt tgtgaacttt cattgatatg tggtttatgg 300
cctctggtaa tcgattacca agggtgggta atcgattaca aggcttataa atgaagacaa 360
gaggcttaaga tggtctctgg taatcgatta ccacggggtg taatcgatta 410

| | |
|-------|-------------|
| <210> | 449 |
| <211> | 455 |
| <212> | DNA |
| <213> | Glycine max |

<223> unsure at all n locations
<400> 449

tagaagaaca aaattgcctc aatcatttcc aaatatgcat gtgaattang aagcatcaac 60
aagaatcaag ccaaggctat tgtgcaagca atcaatgggg caaaacacac caaatgatta 120
tgtatgtatgga tggctcaaata ttcacaaag gtaaactcat cactttcaaa ttgagcttc 180

aaaactatca ttacatgtag aggagaatca aggatttcaa gtcacaaaaat gtcaagaact 240
tttattttca aaacaattac ccatttcctg aacatatccc ataattcaaa gaaaaacatg 300
caaagtgcata catgcacaca aaattgaccc aaaatattaa actaaaaatc cgacgaaact 360
aacaatatta acaaattaac acaactaaca aattaacaaa accaaccaaa ctagcanaac 420
caaagaacac tccccccccc atacttaaac aatac 455

<210> 450
<211> 211
<212> DNA
<213> Glycine max

<400> 450

ctacatgttt atgtatgagt gtgtgcattgc ttgccagcat attcatagat atctattaaa 60
aaaataaatt aaattgaaaa caaataaatt taaagcgtat tgttatatta aaacaaatat 120
gtttaaaacg catatttata ctaaaattac atgatttga atgtttatat ttatactaca 180
attatattag agcaacttaa atttgtgtat a 211

<210> 451
<211> 384
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 451

taatcaagat aagtatgaaa agggttnttc ataaactaag tagcacatgg attnttctca 60
naacatgttt accaaagagt ttttactctc tgtaatcga ttaccagatt attgtaatcg 120
attaccagta gcaaatgga tttgaaaaag tttcaaact gaatttacaa cgttccaatt 180
tatttcaaaa agctgtaatc gattacaatg tttggtaat agattaccag tgccttgaa 240
tggtgaaatt caaattcaaa tgtgaagagt cacatcctt cacataaaag ctttgagtaa 300
tcgattacat tgatttgta atcgattacc agtgattgtt tctaaataaa tcaaaagatg 360
taactcttca aatggggttt tgac 384

<210> 452
<211> 401
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 452

tggatttcct tntagtaggg aatctatcct tcctaagatg gagccaaacc cagtcacccct 60
catthaagaac tagcttttt ctcccttat tgccttagt tgaatacacc tttgttggt 120
tctctatttt gttcttaacc ctctcatgca acttctttac aaatttgac ctagattccc 180
cttctttatg tataaaagaa gtgtccagtg ggagggaat gaggtctaac agtgttaggg 240
gattgaaccc atagacaacc tcaaaaagggg actgcttggt ggttctatga accccccctgt 300
tgtaggcaaa ttctacatga ggaagatact catcccaaga cttatggttgc cctttcagaa 360
gagcccttat aatggtgat taaaacctat tcactacctc t 401

<210> 453
<211> 444
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 453

tcgtgcatgc ccttctccat ttcttgatc agtgacttga gttcgctgtt tgaacggccc 60
aacgcattgt tggccgttt catctttcg attttcgag ctattgttt caggatattt 120
tggaggcaga ccaagtttg tttaaagagt gtggctctgc ttattgttcc taaaacaacc 180
aattctacga gcatacgctc tgccatatta attcacgagc tagctagtct caataacgag 240
aagatgtata tatatagcaa ctttagactc tgccacaatat aaactcatgc atgcaaccaa 300
gatgcttctt gtatactatt gttaaaggaa agtttcttg acacttcatt cccaatttca 360
tattgcttct tatantattt gctgacttgc actggtagc gttaatatca acgggtgcaa 420
aacgtctttc ttttcttttgc acct 444

<210> 454
<211> 453
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 454

tcttccttaa atcaaagtat aaatgattnt atattataag tataaaccta taaaaaaattt 60

ccgttcaaaa taataagata atttacactg aaatttgaa ccactccact tgttttaat 120
gagccaattc caatcttctg gagatacgta tttggctca tcattttctt taagataagc 180
ttgatgtaaa tgataaagct tattcttgc ttgagattga atttttca ttgcatttc 240
tcttcctcg ccagataaac taaaagcatc cttatatcaa acgtttcac cagcaaaagt 300
aaaaattaaa tactatcaag atacattatc aaaccaattt gaagatataa ctaatattca 360
aatatataaa taaatttagat ataattaacc taatttatata ctaacattca tatgtcgcca 420
catttcatct acaccatcct tactaatttc act 453

<210> 455
<211> 442
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 455

tgttcttagt aagatcgatc agacgatgct tttgttcctt ggagagcaac ccgtcttggc 60
cccagaaagt gttgccaatg ctcttagtgg agttagtggac gatccttct cacacttaat 120
cgcctgttga taaaattctg caccaaatat ccaaacacca taaaataaac taaaattcaa 180
ataattaatt gtatggatgg attagtcaaa taaaaatttc tattagccaa gatattataag 240
gaatttgtca agtctatcac gtgagcagtg ggcttagtgt tccttcatt aggagattca 300
tgaaaaattnt gtcggccaca naattgtgag acttcttaat caattaaatt aaaaaaaaaacc 360
tagtaaaata aaagttgacc agctanact aatagtggga atataattag tatttgaatg 420
ccaaaatcaa ccacatattt tt 442

<210> 456
<211> 431
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 456

cgagaaagcc ctctgattc tgtttataca tttctgactc tatggcatga gatgaatgc 60
acagattggc ctcctgtta gttgttatca aagaatagct tatacacttgc tgcttgatgt 120
aaacagttgc tgtgagactg tggttgagc tacttcctt gataacctgtc ttatgattaa 180

cttcatctaa ctgtatagtt cacattttgt ttcctcttt gtctagctgc atattctggg 240
aaaacaagtg ataggtacac attgcttcat ctttacatc atgcaatcaa taaattntaa 300
tgcatacaccc ttgaacata aacactgcat gttntaccac ttgaggacaa gtgagttgtt 360
ctctttgct tgaggacaag caaaaactatt aaatttgggg agttttagt cgatgaatac 420
gactaacttt t 431

<210> 457
<211> 357
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 457

cttctcgat atctgttatt gataaacaag atcaagatga gtatagtttgg aggtttatac 60
acagaatctc caagattgtt tatagtacaa gcggtagta aacacagcat atatgatacg 120
tggcaagcac atggtagag gtgaatcggt aatcagtaat cacacagtta agtggaaatac 180
taaagtttat ggagatagtc cgaaatctt cagaaagtca aaagttctt taatagtagt 240
taaataaata aataaataaaa agtagaattt gactaagaat gaaaaagtaa tattttgttc 300
aataactcat ggnggtatat atacgagtgt tctgacgcgg gaatcacaaa cagaacc 357

<210> 458
<211> 479
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 458

ctattaggac tcttgcaact cagctcgat atatagtttgc aatccgaggt cttcaagag 60
accttagtaaa agtacgtgcc tattgttcat tggaaaccaac aaagtcaatc atgattccc 120
ctgacaacac ctttactctt gcaaagtgc aatctatcta tgtgtgttttgc gtttgcgttcat 180
ggaagaccat attagatgca atgtgaagag cggtttgatt gtcacaaata agcttagtgt 240
cctgagtgcc taagtcttgc aatttcgcgttcat gtatgtgcata ccatggcaca atgttaagct 300
ttgatgttgc acatgttgc ttttttttttgc tccatgttgcata caaatccct 360
caaagcaaaa cacaataacc agaggttagat ctccccgttca tggntatcga attgaaagat 420

tcgaattgtg aatcagaaaa gctatattac gaatcgtaa tcgaatcata tatgaatcg 479

<210> 459
<211> 456
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 459

tgtcaaataa atcactgcat attgtgcagt ccctatcatc cttagagga atgtagaaga 60
gagtaggctt gcaaacaagg ttccaatttg gataaaagtg taaaactact tattaattag 120
aattttatga atcattgttt ggaatattga agaaaaaaaga caacctgaca accagctgcc 180
ctgaaggaac taaaatcatg gcatccaaca agaactctgc atgcttcctg catggatatc 240
atcaataact tgaatagctg caagatttag aatctagtaa gtataaggag ctgttaacta 300
acttgcatacg ctggaagact aagctcctca ggtacatgcc atgctcgatc tttctcgaag 360
gttgacaaag gctctggccc agaaagcaac cgatagaagt atctgaatag tagttccaaa 420
taactatcaa ttactgcana tggtcataaa caaaac 456

<210> 460
<211> 470
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 460

tcagacccaa gcaactcana atctaggat ctaaaacccc tcaatttagt ggattttcaa 60
ggtttgagaa gtgaaaatga gaatgggtt aacttggagc aaactctcat ctcaaacaag 120
tctatatcat caatctaaac tcgctcaaacc tggttttacg acgaaaactc taccgaatca 180
aaatttgact cctcaacacc caatttacc ctagaaatgg ctcttgaaa cacttggtc 240
actcatattc ctcatttgca cagtctaagc tttctcataa gtcctaaatg acatttcaaa 300
ctaggattaa ctccctttaa cctccaaata ccactaaatc cagattggc cttccaaatc 360
tcaaggctca ctcttttcc actcataaca ccacattctc actntctaac ccttaggttaa 420
ctctaccctt catctcttagc agttgtccat aagcaatttc agcacataaa 470

<210> 461

<211> 248
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 461

 tctacttatg tggcagggcg ggcttnctt actttctgtc tcaacgcgag ctttgaacac 60

 tgttcttcct tcccgcgatg cttctttca tgtccgcctg agtgggctta tagcctaaac 120

 catacttccc acgatttcct tgggtattta tcacgctagt tatgcccca ttgtcttgc 180

 ctaaacccat cccgggttca taaccgttcc ccaacataac tcgggcccattt acccgctg 240

 cgtcggac 248

 <210> 462
 <211> 216
 <212> DNA
 <213> Glycine max

 <400> 462

 taataaaaat attatttaaa atcattaatt gagtattatg aattaatata attgttaaaa 60

 aattatagag tattagaaga caacattgtt ataaaaagcg actctattat attgttagata 120

 aggtcaaagc ttttattgtg aggttctgtat atattgttag gaagttataa atttattacc 180

 tcattaagta tatttactat gaaaaaagtg actcta 216

 <210> 463
 <211> 449
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 463

 tctagccaaa tggacttacc ttgacttaat tccttgata gccctttga gccttgttcc 60

 ccttccttg ttttgaagct cactacaagc cttaagtgaa aaaccatgat atcactatat 120

 ccttaaggaa ttttggagct ttggaaattgt tttggataa agtgtggggg attttgttcc 180

 atggataac atgtttgtt ggccatgctt cattatataat tttgagccat acttgataca 240

 cattgcataat tggtaaatg ttggacatgc tgaatatgat gttgtttctc aaaaggctac 300

 agaaaaaaaaa atattataaa aaaaatcgaa aaagaaaaac agtaaagttg agtgaataag 360

aaaagaatga tgagactctn gggttctactc tnntatgtta aaatntatct ttacttcttt 420

ttattttctt atggtttctt aatatgcac 449

<210> 464

<211> 432

<212> DNA

<213> Glycine max

<400> 464

ttcaggataa ggatgaaaga aggcaagatt gcacgtgtat ttgtctgcat catatgtcat 60

agcttagcgat aaatcgtaga cagtagcgat gaagcggca atgtctgttc agaaaatatgt 120

cgggttggga cagttgggtc tcatttcagg catttgcag gctggcttgc gtctcctatt 180

ctcttcatag gaaataatta attcccaatt agcaaagaag attaattaat tgaatgcttc 240

agaaatttcc ttaatcttg agtcacagct ttattattat taattatatt atttcttctg 300

tctatatatt atatatcgct gtaacgcgta tcatttcattc atcaaggaaat ggtatctctc 360

actgttaata gaaaactacc aacagtacag ttcttatct aacccttga agtgcggagt 420

acagttctta tc 432

<210> 465

<211> 442

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 465

ngaagtactc aagggatttg cattaggtag ttcgagcaat actccttgca gaatatcaga 60

tcttgctgca aattcagaaa gactgaagcg actcagcata catataggaa gtggggctgt 120

gatctaagaa ggagaggttg aaagcttgag agaactgtca tcacttgagc atctcaaaat 180

atcatgggt gtgtcagaca caaggtacgg tgatattcca atcagttgc ctccagagtt 240

aaaaatgttg caacttgaag gctttcctgg aaagaatttt ccagaatggt tgaatattca 300

tagtaagcta tccagaaaaat ttatgtcact atctacgata gggggaaaac ttgaaagtat 360

ggatattctc aaatatgttt accagtacat ggggatccta ngttcaagc atttgattct 420

tgacacacca cnatttgaa ac 442

<210> 466
<211> 347
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 466

tgcttggcaa ctggttactg ctgaatccctc ttgggtcttt ttgataaggg ctagattttt 60
tagagggAAC cacagAGTTG cttccacgt ttcttcttct atatggtata gnttatgccc 120
cttcttagat gtataactg ataactctca atggcaaatt ggTTGGTAA aacattgcgc 180
tttggactga caaatggctc tcccaacccc taagtgcatt tgTTGcatat tcctgaatcc 240
tatcataacca atttaaattc cacagtggca gactatattt ataatggtgt gtagcgcatt 300
cctcaatctt tgtagcaatt atatccgact ttgatgaatg aaattca 347

<210> 467
<211> 415
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 467

ntataagcgc gggctggga gacgaaggc aagtggtcgc gatatacgaa gatgatgttc 60
cgagtacatt ggatttggta cgaccatgcc ctcctaattt ccagctggga aattggcgag 120
tggaggaacg cccggcatt tacgcaacga gcataatgta aaccttacg gtttaaaag 180
ctctatagtt gggcttaggc tttagagttt ttccctttgt taaggcttg tgtctttgt 240
tttgaattta taataacaagg atctttcttc atctgttcct acgtctctac ccattctcat 300
tcatttgcat gttacttct tttctgana atggcagatc cgatgacgag tccccgtaaag 360
gtactaatac ctgggaccccg cctatcgact tcgagcaaga aatgagtcaa acgga 415

<210> 468
<211> 329
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 468

ntacagcaga atttagtaat gacccactaa cctagaatta aatataactt aatgccatta 60

accttagggaa ttaaaaacaaa ctaaatggct gagtgtaact gaaattgttgc acaaccaaaa 120
gtcaccccca acagccaaca agtcagccac catttggtct cccaaaaggc tgatgcctag 180
gttgccaatt gggcccttat tacaacttga actaaagccc ttttagttga ttaacccaaa 240
acatattttt ggtcagccaa cttaacaagg attgggcat tatttagaca aactaaacac 300
tctaaaantg aaataaaagtg gtgtcattt 329

<210> 469
<211> 359
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 469

agtcggcaca ttctggccaa ttgcattgtct ccctttccct ttccctctgc tcattcgatc 60
agctcttaaa tctaagaaca tccaaatggg aacgtgaggg gtgatcagaa cccaaaatggc 120
caatcttagt agtcctggat ttaaccttctt ttgggtgtct ctcttggtta cggtggtcca 180
aatccaaacc aaggtgcaat gctaccagta caaagttgga gatctagatt cctggggat 240
ccccatttca ccaagttcac acctctacga caaatggtcc aaatatcaca acctcaggat 300
cggtgattcg cttcgttaagt cctcccttga tgtccatnn tttattgaga catgttaat 359

<210> 470
<211> 454
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 470

cttggagttt ccaagtgcataattcgcttcc ttcttttagtc cagtcattttt ctggcttcaa 60
ttcatcagtggc ttccatcgatc ggctttccctt ctgtgtccag catcttggga tgccatggc ctttgcgtac 120
agctttccag gttctgttat ccagtgtttt gagaaggcc accatccttgc ctttccagta 180
ttcatagttt gtccttccatca gaatttgtgg tctgttcaact ggtccgcctt ctttctccat 240
gttcatcaga atttatctcc ctagatctca ctcagtgtt tcgagtgccct gctctgatac 300
caattgaaat tctgtatactg nggacagatg tcgtacagga tgcacgaca tcacgcttca 360
gaacatgcag atttatatttgc acagtgttgc caaattaaac aagttaataa cacaagagaa 420

ttgtaaccca gttcggtgaa cctcactaca tctg

454

<210> 471
<211> 426
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 471

ggctntgcan aatatatact tcattcagtg gaaggatgag aaatactaac tctntattct 60
tacaagttag aacatacaag aatgcaatca gatgttcata gaaatatttg ctgtacccccc 120
tacatcctca aatacactat ctctgattca ataaaactag agaatatcat tataaccatc 180
tctgaattga taaggaaaaa ggagttaaga gcttgattaa catcgccctc atttatttagc 240
aatttctcgc aatatcaagg atcacgatga aacccaaatc gcggccacaa tttaaaacct 300
tagcacagat gaaatctcaa gttaaacacat aacaataagc aaacacccatc agcatgatta 360
acatttcaac attcaaagat cctcccatca caacccacat aaaactcata cctcacanaa 420
cacatc 426

<210> 472
<211> 453
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 472

gttgaatgca tttaaggtaa acaaacccaaa agcaagaaat tgtgtgcata tagagctaca 60
gacgtcttgg aattgagaca tacaaacatt tgtggccat ttcatacact ttcatggaat 120
ggtaacaat attttatatc attcatagac gattactcca gatatgcata ctgtttctt 180
atacatgaaa agtcacaatc tttggatgta ttcaaaacat tttaagttga agttgaaaat 240
caactccaca aaagaataaa gtgtgtcaga tctaaccatg gtggtaata ctatggcaga 300
tatgacggtt caggtgaaca acatccggng cctttgccca ggtacccatg ggaatgtgga 360
atcgccccac agtacaccat gtcggngtca cctagcatga atgggtggc tgaaagatga 420
aatagaactc ttaacgatata ggttaagaagc atg 453

<210> 473

<211> 432
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 473

ntgagcaaat tcaaacgaac aataactttt actcgatgt cagattgagt cccgtaatat 60
atcgaaaagc tcgaaattga atgttgaagc tctaagcaaa ttcaaacgac aaaaactttt 120
tactcgatg tctgatttag tcccgtata tatcgaaaag ctcgaatgtg aatgtagaag 180
ctctgagcat attcaaacga caataactttt ttactcgat gtctgattga gtcccgtaat 240
atatcgagat gctcgaaatg gaataccgaa gctctgagca aattcaaaca ataataactt 300
tnactcgga tgtccgattt agtccgtaa tatatctgaa cgctcgat tgaatgtcg 360
agctctgagc aaattcaaacc gacaataaca ttttactcggt atgtctgatt gagtcccgta 420
tatatcttga cg 432

<210> 474
<211> 368
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 474

tccaccggtt gtgattgcga gataatattc gtggagggag aataaggaat cgtatgaaga 60
cagtacaagt ggaggttca atctcttctc cgtctctctg actttggga attctatcg 120
agtagtcgga tgaataattt aaagaatttc tggaaaccgc tagagatgtt gttatcgctg 180
gctgaagaca cgtgagcccg cttagaggtt agggatgagt ttatcgaaa tggattaga 240
atgaacatgt gtanggatcc ttagagaact aaatttggt taatttgcga tggttattga 300
aatataattt ctcttatga ttataaatat aatattaatg gggtctatgt accaatgtg 360
ttctgatg 368

<210> 475
<211> 409
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 475

ntgagccaaa atcctgactc accatanacc ttgaccagg gtgagaatgt caatcttac 60
cctcggaagc gaaaagaata gaaggaaat ttccaatcaa agaaaaggaa agaaggaaga 120
tttccaatca aagagaaagc aaaaaaagaa aagaaggaaa attcccaatc aaagagtggg 180
agaaagccaaa aagaaaagaa agaaaattcc caatcaaaga atgggagaaa gtaaaaaagg 240
aagaagaaga aggaaagata gtcctgatc agggatcgaa agaaaacaga agaaatgtgc 300
agaaaggtct ttggaccgga caatatctga ataatacaga gttgtcacca aatgaacaaa 360
aagaaggaaa gcaaaccacg acctanaatg gtcttctccc tttgattac 409

<210> 476
<211> 434
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 476

tgagtgtcca catgggtgca tgcgtatca attntgtata agtttctaaa tcatcggtgt 60
tatatacgtg tcatggaaat aatgtggggc acttcctttt atccctgaac cgctggccaa 120
agcaaatacc ctgacatacg tcatgtcttg ctttctttaa agccttttg ggacaaaacc 180
tcaatccttc agccctaagc ctgcacccaa ggttagaatt tttaccctta tcctcgaaaa 240
aaagaagaac aggaggatct aaaaaaaaaacg agaggaagaa aaagttcat ttactttaa 300
gttatgaatg tgccttacaa gaaaaataaa aagagaaaat ccccaatcaa agattggagg 360
atagcaaaag aaaaagaaaa agaacaattc ccgatcaaag atcggaaagaa agcataagaa 420
aaatatacag aaag 434

<210> 477
<211> 426
<212> DNA
<213> Glycine max

<400> 477

tgctcgtaa gcctgtgctt tcttcttgag tggttgcgt aagctcggt tgccgcacta 60
agcgctaatac tttctttgtc ttaaaaaatt gtgaaattag gcttagcgag caggcttcc 120
aagcctattc tgcagaaaaa aagattttat gtgttcttgc gctaagagca tggctatcac 180

gcttagctta tgagtaaaat ttcataaggc gcactaagtgc catctgctgc gctaagcgcc 240
caatcttaat tctagattta tttttgctt ttctttgga ataattcttg tctagtcgg 300
gctttgatt cttttgttt tcagatggct tcatgaaaga ggaagacaac agctgcagta 360
ccccaggccg gatatgacat atctagattt acatcccaag aggcatgtga ctgctacaca 420
tataat 426

<210> 478
<211> 514
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 478

atgtctctgg gcccttggac ctgtananc cttttgnnc ctttgaaccc ccgcggctat 60
ttaaaaaaaaac cccgcctga caagaaagca ctgtggagat gtttaccacc tcttatgact 120
ggaaagcggg ttctaatgac tcctctgcgg tctccacata aggcataatag gaagggcagc 180
tcaccaagat gtcttactcg cctgatacga tgaccagatg cccttncact acaaataatca 240
acttttggtg gagcggagag ggaacaacta ccactgagtg gattcacgga cgcccccaaca 300
gacagctgta gaggaggta atatccatta tttggaaagt cacttgacag gtggggaggc 360
ctattcagtc tgggagaact attctttcc caaaccttct cgggtggtc tctggcaca 420
acccactttt gaccatattt gtatgagaac aaagtgcgtg gtcactctt gggttaaccc 480
atgaattgat taatgattgt gaatatatcg agag 514

<210> 479
<211> 456
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 479

ctattacgtg acactatgaa actaagctnt aacanatgtc ttcacaaata atcatcacac 60
agcagaaaacc tagcaagact acccatcata tctcccaaa cccataaccc aaaaaattta 120
aaggagaaaag aagtccaccc aaacctgaat tttcgaagtc ccactcgtag ccacgcactt 180
cacgaccccg aaaatgccct ctttcgcga tttggggcag aaatgatggc caaagggttga 240

agctttgctt ggagcttcaa tggagaatga agaagaagaa aatggcaacg tgagagagaa 300
agagagctt ctgaaaagtg tgggggctga gtgaagagag agaaaagctt tttggttnta 360
aataaatggg gtntctcttt ttctattatt ntatttaagc aaatgccaca tgtctncatt 420
tgagtggagc aaaaagggcc cactttcct tttgac 456

<210> 480
<211> 519
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 480

tcccaccgtt tttgaagca tgctattagg acctatgaaa ctaagctgtg ctanaggaa 60
aacttanata aagataactn tcatttcaca tggtaggtgt agctgggtta ttcggcggct 120
cctgtataag gataaggggc gaaaagtaat caaatagaaa taaaactaag aaggcaaagt 180
gtgaatgtgt taaattgtca ataatctcac ttcaaccatt caacacagat acataagatt 240
atttcagcct aatcgataat ttttaattta aatataagta tataattgtt taaaatattc 300
aaaaagaaaa ttttgtaca caatcatatt acaatcaaat aaaattgtgt ctccggtag 360
agatacatat agtccctact caaaaaatcc catttcaaaa gtaaataagg caaattacta 420
atgttggaaa gtttgtaat catgaaatat gtcgcttct tactagacct gacaggtcat 480
ttaatatatt attatcaagn attgttacg cctatgaga 519

<210> 481
<211> 245
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 481

tgagctgcan atgtgcctc accatctgtc aactgtcacc tgtacaactc tggcacgtgc 60
ccaagtggaaa tgcacatgtg ctttatctgt aactacacat ggcttctta acgtcatcac 120
gagaaagagc atgtacgtgt agggcttccc aattaacaat ggaagatgtt ggaggcatga 180
catataatcc antgtgtgat ggagggagag gggttggaa agaaatgagg ggtgggtggta 240
gggggg 245

<210> 482
<211> 451
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 482

ctgcccattt aagtggcc agcgaanaaa tcgaagtggg tctgagaaga ggcaaattt 60
attatcctgc tttgatgaat aggaagctt gggcaaatgg agagaataag aaggagggaa 120
aatcagtcct ttcattacc caccaccta ccagccatga acgcctaatac atccacaaag 180
gccatcccc aatcagccaa aaatccaccc gatgcacatc caagacccaa taccacccct 240
aataccaatc aaaacaccaa ctagggagg aatttccag aaaagaagcc tgtagaattc 300
accccaattc caatgccata tggtgactta ctcccttacc tgctcgacaa tgcaatggta 360
gttataagcc caacaaaaat ttctcaacct ctgtttccca gaggatacaa ccncaacgtg 420
acatgttctt atcatggggg anngttggg c 451

<210> 483
<211> 449
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 483

tctataaaag gtgcgttcct aatttctcta caattgcattt acctctcaat gagctggta 60
agaagaatgt ggcatttacc tgggtgaaa aacaagagca agccttgct ttgctccaag 120
aaaagcttac taaggcacct gttctagctc ttccctgattt ttctaaaact tttgataata 180
ttagggactt gtatgctta gatgaacatt tctctccat ttacgaaagt tgtggaaaa 240
aggcccaaaa tggattctat ttggcttaagg ggtatttgtt caaagagggaa aagcttgca 300
taccccaagg atccattagg aaattacttg ttaaagatag ccatgagggt gggctcatgg 360
gccactttgg gatagacaag acgctcgatct tactcaaaga anagtttat tggcccccata 420
tgaagaaaaa tgtccttaag cattgcact 449

<210> 484
<211> 405
<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 484

actctcttat cgaatggtgt gccacaataa catcaacatc gatttgatga ttgcattgctc 60
ccttttctga tcttataaggc acgttacagg gacgaaatca ttatttaaat atttaagtgc 120
ccaaagattt tgattntaag agcaaaataa aataaaaaaa atactgaatc atgtggagtt 180
tattgaactt catatgcccg agtagagtgt catgcattt aatttaact gaatcaaact 240
aaactgtacc tttcacatta gtaatgtaat tagtaagact agaagcatct taatgaggtg 300
gcagaatnta attaattatt tagaaacatc ctaatgaggt ggcagaattt aattaattat 360
ttgatttcaa tattttcat actaatttct tctnctttat ccctt 405

<210> 485

<211> 398

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 485

tgaagagagt tttccatgc cagcacgcgc ccaacgtgtc tgttagctgc taagcgagat 60
gggtgtctct tctgcgctaa gcgcgagatt gccgctaagc caaatctcac ttactcgac 120
ttagcgcgag aatggcatta aacgcgcctt catggacagg aagcccttc ttaagcctga 180
cttacagaaa atgaagggaa gggctggaag agagcgctga atagccgtca gagtttgaag 240
agtgaaatac acaaaggcaa ataacagagc anaggagcca agttttgatc ttttaggaag 300
atttgtgagt ctttgagtga ttgtgagatt ccttagaggtg gaggagacat cctcactcct 360
ttttagcaa gcaatntctc ttaattcctc ttctttca 398

<210> 486

<211> 382

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 486

tcattgatca attctgaagt tcacaattgt cataaggatca tgaaaggagc aaaatgttgg 60
atcaagtggc ctcagaataa ttaagaaggg gggttgaatt aattattaat gagcctttac 120

taatcaaaaa cttatccttc ttaatgttac tagattcaat taggcttta ctactaagt 180
aagaaaagtaa agaacagaaa tagaaactta atcaaatgta aaagcaataa ttaaagtgca 240
cagcgaaaat taaagagtat agggaagaac aagacaaacg caagaattnt atactggttc 300
ggcaaaaactc atgcctacat ccaatcccc acaacacctgc gttcttgag atttcttca 360
accttgtaaa atcctttaca ag 382

<210> 487
<211> 399
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 487

nttggcttta gaattaatca tcaaaagtct catttcattt gttctaaagaa tattccgaga 60
aggaaaactc anaaatttgc tatattttgg gtttccaata tatactcatg atattggtaa 120
atattgggt ttacctataa ttagtggaaag agttaaaaaa aaccacttct cgtttattct 180
ggataaagta aatgatcgct gagctggttg gaaatcgaag cttctcaata gacttggc 240
ggttacactt tgcaaatatg tcctcaattt tatttttttttatacatgc aaaacatgtg 300
gctccttcaa ggcattttgttattcccttga tattgctact agacaattca tctgcggatc 360
aacttcatct cattgggtga gttggaaagac tnatcatc 399

<210> 488
<211> 396
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 488

tcttggcaat cctcattcca gcgatcagtt tggttttgc gtaagagttt gaacaacggc 60
tcacaaatgg cggtgagctg cgatatgaat ctggcaatat aattcaagcg tcccaggaaa 120
cctcggactt gcctctctgt acggngttct ggcatctcaa ggatagcctt cacctttcg 180
gggtctacct ctatcccttt ctggcttaca acgaaaccaa gcaatttccc tgatttgacc 240
ccaaaggtagtac acttagcggg gttcaacctt aattgatatt tcttaagcct ttcgaacaac 300
ttccgctgggt tgacaagggtt ttcttcctcg gattttagatt tagcaattat gtcgtccac 360

tagacctcgatcgtatcatatatca tgaaac

396

<210> 489
<211> 383
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 489

tcatggtaa tcaaaggta ttcanaggtt ttttgatgt aacaatgtatg ataacaaaag 60
atgtgacaa aggtgatgac aaaaagctca aagatcaatc aagaacaatt caagagttca 120
agataagaat caagaagaat tcaagactca ataagaaagt ctagagacaa gaatcaagat 180
tcaaggttca agatctcaag aatcaagaac aagattcaag actcaagatt caagaatgaa 240
gagaagactc aatcaagata agtattaaaa agttttcaa aactttgaat agcacatgag 300
ttttgacaa aaactttac cagagtttt actctctggt aatcgattac catatagttg 360
taatcgatta ccagtagcaa aat 383

<210> 490
<211> 346
<212> DNA
<213> Glycine max

<400> 490

tctacttatg tggtagggcg ggcttccttc actttttgt ctccaaacgcg agctttgacc 60
actgttcttc ctccccgca tgctttttt catgtccgcc tgagtggct tatagcctaa 120
accatacttc ccacgatttc cttgggtatt tatcaggcta gttatgccgc cggtgtctt 180
gcctaaaccc atcccggtt cataaccgtt ccccaacata actcgggcca tcattactgc 240
tgcaacggac agacaagggtt gcccagagag ggagtccacg gagaaatgc tgaccaccc 300
ataagactgg aaagcgggtt ctaacgattc ttctgcggct tccaca 346

<210> 491
<211> 391
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 491

tgaaggtaaa aggttacac agttcatgt taaaagttac tgacatcctg ctgctaata 60
acatttagacg tagctntact ctttagtaaga atattagaac aacacagcaa acaaaacact 120
ttctatgcac tgagcaaatg tattcaaaaa ataattatgg attagaacta agtttcaca 180
aatcttaagc aagcatcagt aacatctta cctgcagcac tagaaagaac ccaattgtca 240
tcataaggaa ctgccacac agcatagcca agtaacttat ttcccttgc ataagaaacc 300
ttcattntga caacctaaca atcatcataa cctatccaag tcgatccatt ggagaagtaa 360
ttaactacat aagtagcatt gtacttgaca t 391

<210> 492
<211> 454
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 492

ctcagcttan atatgtttc cctcanatga nnactaattt agttntaacc catcaattaa 60
aaatttgctt tctacaaatc ctcaattca cttgaggca ctgaaatcaa cagtttacga 120
atttgagtga tttaaaacat aaattatgct catgtaacct ttcatgaata aagactaaat 180
tattcattca aaaagaataa atactgaaac agtatgaagt taatgaattc attgactaag 240
taaatataga aggacaaatt aaataaaaaa attatattat atgcaagtaa tctttaatt 300
ttaatataaa attaatttagc acacactatc actttgaaa atgatatata tatatatata 360
tatatatata tatattaatt ctaacattag tattcgtata gtatacacgg attgtaata 420
ataacaagtg aacaacaata acagtaataa taaa 454

<210> 493
<211> 386
<212> DNA
<213> Glycine max

<400> 493

tgcacttcag gacgtgcaat gctggtaaag aattgtggca ctgaactact tgagttgtcc 60
tcaagggtaa cccttggta gtgttcatca tccatggtaa tggcctctgg taagtttgaa 120
gctgattccc ttgataatgg tgagtcagat gatgaagaat cagaaaaatg agtttcttcc 180

tcaagaatgg atgctactgt cctgtcttgc agtaatttc ttctccttt ggctctgttc 240
cttcttaatg tagcttcaat ctccaagtcc aaaggaacta aattgcctgt gggagatcta 300.
tgcatttaaa acactaacag aaacaacagt tatccagttc aagaggaaaa aaaatatgaa 360
ttaaaagcaa atattcacag ttaatc 386

<210> 494
<211> 443
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 494

tagccgaatt catattgaat tgaagttgc ttagctcaac ctggccagc ttagtgacc 60
aaatcaacct tatgcaaggg ttggcgctca agcacttgag actcacaact tagcgcata 120
accgagatgc gcttagcgta aggcttgcgc ttagcgaaag gactatTTT cagagaaaag 180
ttttctgtta ttttcagtc ctTTTccaa gaaattgaaa cctttatgtt aaacattcaa 240
agataggtt atatactcct atgtacagat ccgacagcaa gttccaaatg attaaatgca 300
tgaaaaacaa agataacaaa attaaaaact gggttgcctc ccaggaagcg cttcttaac 360
gttattagct tgacgctntt accttactgg atgatcttat gtttggttc ttactttcag 420
aacctcttga cctccttcca tta 443

<210> 495
<211> 369
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 495

nttagcttct ctaggaatct tctcaagggaa gtttctcaag gaagcttctt aatgaggtga 60
gcttagttat taggtgtgtg tagctaagtc tagcttctca aggaagcttc tcaaagaagc 120
ttctcaagga agtttcttaa ggaagcttct caaggaagtt tctcatggaa gtttctcaag 180
aaatTTTCTC aaggatgcta cctaggctat aaataaaagc atgtgtaaaca cttgttgcaa 240
ctttgatgaa tgagagtctt gtgagacaca cttcaaagtt caacttctct ccccttttta 300
caccttcaat ttcatgctcc cccctctctc tttctctctc tctttctttt tctccattga 360

agtttccta

369

<210> 496
<211> 372
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 496

nggatgaagt tegaaatctc ttcccttcttc atctctaaat ttcatccacc accacttgca 60
tcaccaagga gagctccaaa actgtgcattt cctcaagctc ttactctact ttctcttaaa 120
ctctctacta tgagtgttt taagtgtgtt aaacccaaat aatccttgcgt gtatTTATAG 180
ggtaaatgtg aggcataagg agtaaataag accaatgagt gttaggatc atataggct 240
tttagttaca aattaattgt tcttatcttt taattntatt tttttctttt cttttattaa 300
tttagatattc tagatgcttc atgggttatt agagtagaga ttgaaatgta tgtatacatg 360
atTTTgatga tg 372

<210> 497
<211> 387
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 497

tttcgcanag cttacggtaa aatctaggac atagccatgg caaaggctta cacagaggcc 60
attgcctccc tcgcccagta ttatgactag ctgttgaggt gcttcacctt tggggacttc 120
cagctatcac ccatggtgaa agaatttgaa gagatcctag gatccctct atgggaaagg 180
aagccatacc tcttctcagg attctatccc tcttagcta gaatttctaa gatagtccaa 240
atctcggtgtt aggaatttgcg ccacagaaag caagtcgaaa atgggggttggt tggagtaccg 300
agaaaatgtt tagaggcaaa agcaagaatc ttggcaggta gaggcaaatg ggccccgttc 360
atagacatcc tcgcactttt gatcttc 387

<210> 498
<211> 446
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 498

tcagcatctt aaacagaggc taaaaatctg aagtaaagac aacatatgag acctgtgcag 60
catagtcaaa cagttcaat agaaaactaaa tgacttggag aattctatgc cagttcaacc 120
ttctgaacaa caagtgaagg acctcaagaa aacccaagct gacctttggg aaaaagctac 180
tatgcaggag tctattgtga ggcagaaatc aagatgtaga cgatcatag agggggacag 240
caacacatcc tatttcata gagttattaa tttgaggagg aagagaaaatg ctctgagggg 300
gttgcagatt ggtgacacct gtgtggaaaa tcctaacatt atanaagctg aaacctttca 360
tcattttaga acaggttcaa tgagcctcac ttgaccagac ctaacttgga tgggggtttc 420
attnaaagtc tgacttattc tcacag 446

<210> 499
<211> 402
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 499

tcaacatgaa gcaatcagct cgcctggca agcatgtgc ttctaaacta agccaccagg 60
ttgccgggtg agctggcg 299 120
caagatcctc ccctatgg gctataaaag ggtgtggag
gctaaggaga aggggttcag cacccttgtt cacttgcagc agacaaggaa agtgcgttg 180
acactgtatg caaaaagtac gacattggca atgagaagtg ggctcaattt tgtcagaccc 240
gcagagaccc ttctgtggag gcaacgtttt tatttcatt gttntaaact ctaaattcac 300
tttagtataat acattgtaat gataacttac aataatggtt aactttaca ggatatgcga 360
aaaaaggcat aagccatcta aaaacaaaac actgtccctc ac 402

<210> 500
<211> 449
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 500

tgcttgtggn gcttctatgg aggctggatc tttgagcttc aatgagggtcc ttcaatggtg 60
attttcgacc atggagatgt agcagaaggc aaaggagaag aggagagagg aggcaccatc 120

cacaagggaa taagccaagg aagaaggagc ttgcacca agatgagcat tggataagaa 180
gcttggagat gatgcttcaa tggaggaaaa gaaagaggga gagaaagaga gagtgaggag 240
cacgaaattg aaggaagaaa aaggagaga agttgaactt tgagttgtt ttcacaagac 300
tctcatccat caaagttaca acaagttta cacatgttc tatatataga ctangtagct 360
tccttgagaa gctntcttga gaaaacttcc ttgagaagct tcttgagaa aacttccttg 420
agaagctaga gcttagctac acacacccc 449

<210> 501
<211> 373
<212> DNA
<213> Glycine max

<400> 501

tgagtttaac gatgacaaat attcaagaag caaacattaa gttatccgaa ccaagctatc 60
tagtattgat tgctttaggt gttcttata acatacgact aaacttataa gaatggggaa 120
gatgtggagc agtagcctat ggttagaaatt ttccagacat agattctgcg taatcaaaag 180
caaaaactact agaaaagagag tggcctaatt ttagatggaa aggaacacac gttttatgag 240
tgaaaactgaa aatataacaa gcttatatatg atcagaagca ttagttttagg tatctttgg 300
aaaggacat atattagaag ctacgcatga ctaaggatc atccaaagta ttctaattca 360
aaagtcatga gaa 373

<210> 502
<211> 320
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 502

ntngagatca aactntacca ctggtaatcg attacaggaa actggtaatc gattaccaga 60
gagtaaatac tctggtaact tagaaaaatt tgaaaaact tttcttgtaa aacaaaattg 120
tgctatgttt ggaaaaattt caataacttcc cttgcgaagt ctgacttgg 180
tgcttttgg tttttctct tgaatcttga atcttcttga tgactttct ttaatcttga 240
tcttgaacctt gttgactcaa tcacgacatt attctttgg cattttgaa atcatcaaaa 300

ctacttgaat tattcttgat

320

<210> 503

<211> 430

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 503

tgggtggtaa aaananaaaat ataagaaaaa ggtgtatgt tatTTTggga ggagccctgc 60

cgattcctaa atcctcttt cagaggaaaa cgttaaagaa atgctactgg aataagataa 120

gataaaaaat ggagatgaga gcatgctcca ctgatttagat tctgtaagcg ttagctacta 180

aacttcatca acaacatggt aagtaattat ggcaacagtg atcgaaccaa aggaatatgc 240

agacaagata gatagagaga aaggaaaaa aagtcaagtt aagtgagtgt catcggtgaa 300

aatcaatata gatgccaggg acggctctgt aaattcagta gcccaccta caaataagtg 360

ccctaattat ctgaacactc ttctcaaata atacattcac aaattaaaca gttccattag 420

taaaaatatg 430

<210> 504

<211> 463

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 504

tattggccgt tggatgaaac tccacacagt gcagttctat gttcacttgt agaggatctg 60

aatcccagtt taccagttgg ttgttgcgt taaccaacaa atcttatttt atgttctctn 120

taatatttct ttntaattga agatgcttct tttacccaca gaaattgtat attttcatg 180

caagtataac agttaaatga atagtagtta gtttaattgt atttcgaatt tttttttata 240

attaaattat aagtgatttt ttagttctta tggaaaaaat taagttggtt aagttttta 300

ttttaaaaat taactaaatt tattcattnt gtcaaattnt tactaaaaaa ttntcaaatt 360

tttaatttta tctacatatt taggagttta tgaataaaatt gagttagaaa ataattntaa 420

aaattatang taactttattt agaaaaatta gtaaatatta atg 463

<210> 505

<211> 406
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 505

tcttcaccc ttgctcagac tggttaattga acttattctt aaattnntgc ccttggtcca 60
aaatttatggg tgaattttca tttagattatt ttttcgcttc acataaaaatg attctttgat 120
catgtgagaa gagaatgaag cctcagtttag gacagttgat taaattaaga atagactaat 180
aattacaggt agaggaagac caagaaagac tttggattct gctatttatta gaattggttt 240
agatttaaat ggcttctatg aaaaatgaaa aattgttttt taacagaata caatggcatt 300
gtttgattga tataggttgt agatgacctc agtgggaaaa aaaactntgg ttatcactgc 360
tatgtataaa tcattgatat tatttggtag caactcttat ggtaaa 406

<210> 506
<211> 460
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 506

tagcccccaa cgaaagatga accgtgtcaa gttgtacact atgtggagat cattgttagga 60
gtagaagggt ccgaggagtt agagttggac ctgagaacca atgacgacaa tcgggttaaa 120
ccaattgaag aaacatgtaa ctttcagctc gacattaaag aggaataggt tacttgactt 180
gggaaccaac tctcaatgga atataagaat gacttacaac aaattatctg agcacatgcc 240
gacctgtttg catggttcga ggtcaacatg ctaggcatac atccgacctt ccattgccat 300
aaattatcca taatuttaaga tgccaaatta tatctcagag gaagagaaaatgngaaag 360
aaaggtgtta ggttagtgcgg caagaggttt ccaaactggt ggcctaatta tcanagaggt 420
agagtacacc acatggctat canatctatt cctagtaaaa 460

<210> 507
<211> 459
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 507

ggcagtanaa aacaacattt ttttcaattt ttatgtcaat tctctgtaat atacaaagtt 60
atcattggtt catttgatgt tggtaattt gaagtagaat aaacacaact acagatgtta 120
ttttactttt taatttatata tactagattt tcaagacatt caagcttca atgccgtcat 180
ttcctctgca cctactcctt cgatttcaca acagtgtcag tagaaatgat gctaaacccg 240
gctcgatattt ccttcctta tgatctttt ccccaacaat ccattctcat actgagggtg 300
aaaaatatcc ttaacagcat gcccatgatt ttattttcc tgcaagtttgc tcacaacatt 360
aacaaagtag taccttatta ttcattatgg tggcctacc taattcagta tggaaatcaac 420
catagagata gatagtgaag tggaaaaca gcaagataa 459

<210> 508
<211> 413
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 508

tgatatctcg gccaatctt tcagttaaa tactccaaat cattattgtt tctaccttct 60
agtatatttc ttgttggtag tgtattggaa tctgaattct cagttcaca atgactaaac 120
ttttttctt tattgacaga gcaatgagga tattaaccac acgcttgtct gtgccaagct 180
tgaactgata gcagcataaa tggaagccaa tgctaagata aagaaatatg aggagacagt 240
taaacacttg tataaacttt taaagaaagt ttgccagggaa agagatgaag caagagacca 300
gcttcaatttg ataaggaatt tccaaggcatc tactccagct gagacaagca gtactggc 360
acaagttgat catcatgcat gcctncaata ccaaacaag ccatcattga ata 413

<210> 509
<211> 363
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 509

tattggaatc agccaattgc tcagtgactg ttcttaattt tgcattttt aaggtgtata 60
gttggcattt caagcataac cgatttgcaa gagacttgtt catatacaag gattccagtt 120
tcaaccacat tgaggttgggt gtccttctc ttgcaacttc tcttatacgat ttatccaa 180

ggcatagtat gattgcactt ctggctctat ccatcatttc ttatttctcc tttgaactta 240
gagattcaga catcctttct tcttctttaa gagcttctac acagccatgc tgaatcaaga 300
ttgcttnat cttgatcctc cataaccga agtcatttc cctgagaact ctcatatcat 360
act 363

<210> 510
<211> 418
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 510

tgattaccct ggtgacactn tacttccagg aatgaaaatt ggagtgaact tcaaaaccgg 60
tcaacatcgat gctctaagat catggagatc ctttacggat cctactccag gtaatttttc 120
acttgggttt gatactcgtg gccttcctca attagttatt acaaatgaga atactaatag 180
taatgacata gcttatagac cagggtcatg gaatggtctt agtacacgg ngcttcctgg 240
agaaataact gaccaattaa caaaatccct ttttggtagt aatcaagatg aggtcttcta 300
tgagattcag ctcttgaata gttcaactaa actcatgaga agcagaatgc ttccagaagg 360
gtatcaagta cgaaaaat ggtcagatga aaaaaaatat gggattctca aattccta 418

<210> 511
<211> 435
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 511

tagaagcaaa tgctggcatg gtatcgaga ttctgttagc aagaccacct gctgaaggac 60
ggngaaaggaa tcaattgctt ccacgttatt ggcttaggat tactgatcaa gagttgcaac 120
aaatatcagg agagtatcca acattgtaat ctgtcctata aattattatg aatcactgac 180
aggataactta ctgggtgata cacttattat tattttgata gctaattttt accttagtat 240
ttccttaacc atgattcatg atatgttcaa gttcaaattc tacaatcgtg ccactcttg 300
aaaagatgct tagtgcaagt gatgctggtc gaattggtcg cttgggttta ccaaaagcat 360
gtgctgaagt aatttatctt aactcatctg ttgaactggc atttactgtt gtcattttat 420

attaactaac aattc

435

<210> 512
<211> 395
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 512

tgtgaatnta tagtgtttgg tgactaattg tcacaaaaaa gcaaagtaaa gcccaaagaa 60
gcaaaattaa agatctaaaa ttactcgctc agcatttctc aggtgctcag cgcaacgcag 120
atgcttagcg gacaacgcac gcttaacgcc agaaagtatg aagacgtctg aatcatgaat 180
atgtgcttag cgcgagtcac tcgctaagcg cgagattact atcatactcg ctaaggcatga 240
aattgcacctt agcgtgaagg ttacgtaaaa atcaaactga actacaccta taaaagaagg 300
agagagaaaa agaaaaaaaa tacactaaa attcaagaga atacaattcc ttacagaagg 360
caaaggtcga aagcaggaga agcaaccatt cgtag 395

<210> 513
<211> 412
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 513

tggtaaagaa aatgatggca tgacttttat ccaatcttat tatgttgaaa agctattgaa 60
gaagtttaat tattttgatg cgaaacatgc tcttactcct tatgactcat ccatcaagtt 120
aaagaaaaat ttgagtaaat gaatttcttt acataaatat tctcaaagta tcggttcttt 180
gttgcatttg acaaacttct ctatgcctgt ctgatattgc acatgcagtt ggttagattgg 240
aaagtaattt agggatttag tggatataaaa ttgaagttct gatntgatg aaataaaaat 300
gagaagtggc tatgtctttg ctttagctag ttgtgcagta tcatgaaaat ctactagaca 360
agttattatt tcacatgana gcaaaaatta ttgctttaaa tactgctact ag 412

<210> 514
<211> 320
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 514

cttgtgtggg acacccattt tgagtgttgt ttccaaaccc ttatagaaaa gttgacgatg 60
actcctgtgt tagtttgct taacccaaga gaaccctttg aggtgtattt tgatgcata 120
aagatgggtt taggaggagt gttgatgcaa aatggacaag tagtggtta tgcttctaga 180
caactcaaga ctcatgagag gaattatcct acccatgatc tggagtttc tgctgttagt 240
tttgctctta acgcgtggag gcattaccta ttcgactcca gtttgaagtg ttttagcgatc 300
ataaaagcct taagtattt 320

<210> 515
<211> 371
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 515

taggaaccca nacttgttagc ttcaatgcaa ggaaacgtgc ttatggctag gaatccaaaa 60
atttggttt agaatttagaa aagaatgaaa atagggactt gttttaaga atttgggctg 120
ccccatgatt ggtactttgc acctaaataa catggaaat gatttcaat gctgtgtaga 180
tatatgtgtta aatatgaagg gcatgaaatt ctttgc当地 gatgaaggaa tattgaggc 240
acttcctaaa tgaatgtatg atagcatggg attccctttt gaatgcaagt atgtgcataa 300
tgttaaatat cttgccaata ggcataagtg tgagtgaaac aatgaaaagt tgtatggat 360
atatatctt 371

<210> 516
<211> 455
<212> DNA
<213> Glycine max

<400> 516

tagcccccta acgaaagatg aaccgtgtca agttgtacac tatgtggaga tcattgtagg 60
agtagaaaggg tccgaggagt tagagttgga cctgagaacc aatgacgaca atcgggttaa 120
accaattgaa gaaacatgta actttcagct cgacattaaa gaggaatagg ttacttgact 180
tgggaaccaa ctctcaatgg aatataagaa tgacttacaa caaattatct gaggcacatgc 240

cgacctgttt gcatggttcg aggtcaacat gctaggcata gatccgacct tccattgcca 300
taaattatcc ataatttaag atgccaaatt atatctcaga ggaagagaaa gattgggaaa 360
gaaagggttt aggtgtgcg gcaagagggtt tcaaactgggt ggctaattat caaagagtag 420
agtcaccaca tggtatcaat ctattctaga aaaaa 455

<210> 517
<211> 461
<212> DNA
<213> Glycine max

<400> 517

tgcttggtt gcttctatgg aggctggatc ttgtatcttc aatgaggtcc ttatggtg 60
atttccacc atggagatgc agcggaaagac aaaggagaag aggtaagagg cggcgccatc 120
caatatggaa taagccttgg aagaaggagc ttaccacca agatgagcct tggataagaa 180
gcttggagag gatgcttcaa tggagagaaa gagagggggg ggagcacgaa attgaaggaa 240
gaaaaaggaa gagaagttga actttgagtt gtgtctcaca agactctcat tcatcaaagt 300
tacaacaagt gttacacatg cttctattta tagacttagt agcttccttg agaagcttc 360
ttgagaaaac ttcccttggaa agcttccttg agaaaacttc cttgagaagc tagagcttag 420
ctacacacac ccctctaata actaagctca cttcttgat a 461

<210> 518
<211> 460
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 518

tggagaggat gttcaatgg agganaagan naaaggagag aaagagagag gggggaggac 60
gaaatttgaag gaagaaaaag ggagagaagt tgaacttga gttgtgtc acaagactct 120
cattcatcaa agttacaaca agtgttacac atgcttctat ttatagacta ggtagcttc 180
ttgagaagct ttcttgagaa aacttccttg agaagcttct ttgagaaaac ttcccttggaa 240
agcttccttg agaaaacttc cttgagaagc tagagcttag ctacacacac ccctctcata 300
actaagctca cttcccttgag aagcttcctt aagaagattc tttaaagaagc tagagcttag 360

ctacacatac ctctctaata gctaagctca ctccttgag atgagaagct agagcttagc 420
tacacacccc gtataatagc taagctcaca tgaaaataac 460

<210> 519
<211> 405
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 519

ntgagccaaa atcctgactc accataaaacc ttgacccagg gtgagaatgt caatccttac 60
cctcgaaagc aaaaaaagaa tagagggaa atttccaatc aaagaaaaag agaaggaaaa 120
tttccaatga aagcaaaaaa agaaaagaag gaaaattccc caatcaaaga gtgggagaaa 180
gcaaaaaaag aaaagaagga aaattccccca atcaaagagt gggagaaagc aaaaagaaaa 240
gaaaggaaaa ttcccaatca aagaatgaga gaaagtaaaa aaggaagaag aagaaggaaa 300
gaaagctcct gatcagggat cgaagggaaa acagaagaaa tgtgcagaga ggtctttgga 360
ccggacaata tatgaacaat acagaattgt caccaaatga acaaa 405

<210> 520
<211> 452
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 520

acaatttatct aatcattcca atccactcaa atcatacagt tgctcattca aatcattctc 60
aaacactcat ttcatacaaa acaatccact gcataatcggtt ttcaatcagt tctctgttca 120
aacaagcttt tttgtacatg caaacaactc atagtactaa aatttaaaga acggaaacat 180
aaaaactgaa atttaaatga ctgaacataa atcataaaat aattcaagta aactaaaatg 240
ttcaaaatgc acaaatttaa atgtcctgct tctgtgggtt ctcttgca tgcttattaa 300
gatccaacac ctgagcaact ggtaaatcct gagaggtagg tttctctaac tcacatgttg 360
gtgaagatgg tatggcatca tcaggtatag gtgctgggaa tggctctggg atctggtctg 420
tggaagtctn cttctttga gccatgtgta ca 452

<210> 521

<211> 391
<212> DNA
<213> Glycine max

<400> 521

tcattctcta tcttgagact cttgtgttat taattactgt ataaacctta gggtttctc 60
attcctatct tctgcaaatt ctccatacaag gctagaaaata tttctgagca aatataaccag 120
atttgattt tgattctttt agcatgcaac tctatattaa tgctgctgca ttgtaatgat 180
cacatttgca acttactaaa atgcggcgca gcttaaggcat ttgaatgctt tttgaatcac 240
tgtatttggc gtatttggtt tggccgttag ccaattcaat gcgtggatat atatcgacaca 300
cctacaccgg agatataatat cagttatggc tttgtgttca agtttctca atcttcatct 360
tctggttttt gtttggttga cgatagaagc g 391

<210> 522
<211> 418
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 522

tcatttgc当地 ttcagtgtaa tcagtgccag cc当地attcga atgaatttggaa tactttgaac 60
cagtttatgt acaaaaagaaaa aaaaaagttt aattttatgt tatcaagtat acgaaatttt 120
ataccatcaa ccaattaaaaa gttatataata atttggaaataa ctattataat aattatcata 180
aaaatcaata aatttatcat tc当地atattt caataccaaat agccactata taattactgc 240
atgctcaatg agaattaaat tgacatgtac ctctgatcgt atagtgtgtt tgaatctgcc 300
atctcttctg atttgaagtag tgcttaaccac gttaaggccct tctttaagca gaatgtccaa 360
caacttgc当地 atggaaaaca aactttccc anagctgtaa ctgcacataa tctcaagt 418

<210> 523
<211> 393
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 523

tcaggctgtt caattgctnt aaattgttgc atagaaggc aaaggtctgt gtgggtggc 60

gcagaggagc acaaaccaca gagtctggcg acaggtgcag atttttatt catggccagt 120
tgggttacca ggttaaccaa tgcatactagt ttaccttcaa gcttcttagt ctcacctgat 180
gaattcgtgg ctacttcatg cactcctcta atgacaatag catcaattct gcactaaat 240
gtgtgggagt ttgaagccat cttcttaatt aaatttcttg ctgcagcagg ggtcatgtct 300
ccaagggctc caccactagc agcatctatc atacttctct ccatgttgct gagtccttca 360
taaaaatatt ggagaagaag ctgctttgaa atc 393

<210> 524
<211> 392
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 524

tctaaactnt aaacaaaaat gaagctaaac tntaaacaag aatgaagctt cgataccact 60
tgttagacaa gtggcctcag atatcttaag aaggggggtt gaattaagat attacaaact 120
atttcccaa ttaaaattct actttgattt taatgcaagt tcaaagttcc cttaaagatt 180
aatttctaaa tgatgattca aaataaccaa actgaatgta aaagtaaagc aacaataaat 240
aaaagagttt aagggaaagag agagtgc当地 ctcagttta tactggttcg gccacaccct 300
tgtgcctacg tccagtc当地 aagcaacccca ctggagagtt ccactaactt gcaaaaaccc 360
tttacaagtt ctgaaccaca caaggacaac cc 392

<210> 525
<211> 209
<212> DNA
<213> Glycine max

<400> 525

ggagaatttg taagacttaa ttcaccctc tcttaagtta ttgaggtcac ttggcgcgca 60
cacagatcaa gaataaaagct aagtcttact ctatcttgc taaaagagtc tcttagtgat 120
tggaaagaat tggcctcaca acattttct taaattgatt ataaaaaggt tctagaatac 180
ttttaacaat ttttggataa gacatttt 209

<210> 526
<211> 431

<212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 526

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tgagcatagc agctntgtta attaccttgt gagagtgaac accaaggcca cccaaatccg 60
taggcaaaca aaccttttc caaaaaacaa taacaagttt tgatcagtt tcccttccc 120
ctgtccagat gaagtttga atccaagagt caattatgtt caataatgta atcggccatg 180
cataaatgtg aaacgaataa agcaacaagc tctaaataac aaatttgatg agaaagctat 240
tgtgtgatca acgctttac aaacaaaatt atccacactc acaacggatc gtgagaatac 300
aacaaagatt ataccataga aaaataataa caaataanaa tttaacatga ttgcacacat 360
cttgccata tctacggagc tggttcaaaa atattgttt atcatataaa taattacaag 420
aatggaattc a 431
  
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<210> 527
 <211> 330
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 527

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tgttgcagac aaaaatctnt ggctagaatg ccgcanaaat ggtctacaaa acatccatcg 60
gttttcttgg cctgaacact gccgcaacta cctctcccat gttgagtatg gcaggaaccg 120
ccactccaca tcccgcttgg aaattacacc aatgactgaa gaatcaataa gtgactcatt 180
gagagatgta gaagacattt cttagatt ctccacagaa ggagactcca agcagaatgg 240
agagatggac actgcagcaa ggcagaagca aattatggaa gcaatcatgt gcagggtctc 300
ttccactggc aagtctaattt ccagttactt 330
  
```

<210> 528
 <211> 450
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 528

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tatcaaacca tatacaagga tcaggatccc ttttatatct aaagtaatta tggccact 60
  
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gccatcgAAC aaattaaagg aatacccaag aaaaaattgt agagcttaag ttgatgttac 120
tttttcattt gataataaga tttattgtt aaattgaata gatacttatg gttaatctaa 180
tactctttat ataacacaaa acccatcaat ttgcaggaac ttaatgttcc tgagcatgat 240
gttgagcagc tattgggtgc actgatttg gataatagaa tccaaggca tattgatcaa 300
gtgaaccggc tcttagaacg ctctgatagg tcttgccgtt atatttgat ttgttaaatt 360
aaattcgtca tattcatctt tctttatat aaaacataat atntactaac atattcacgg 420
tccaggtcga aaggaatgaa gaagtacact 450

<210> 529
<211> 416
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 529

aggatagaca aacagcgcta gnccaatcaa ttgtgggct ccaaactcga tggggagga 60
tgcatgaatg acaagcaatt catgggctc cgataagat ttgaaggttag agatagatg 120
aacagcacta ggcaatcaat tcgtgggct ccagacttga tggatgatga tgcatgaatg 180
acaagaaatt catgggctt tggataatat ttgagggtagg agatagacg aacagcgcta 240
ggcaatcaat tcgtgggct ccagactcga tggggagga tgcatgaatg acaagcaatt 300
catggngctc cgataagat ttgttggcag gactgaatgg tccaccgggtt ttttcccac 360
cctaaaggcg aacatgtttt atcaaggaan aataaatcat tcatgagagc actata 416

<210> 530
<211> 221
<212> DNA
<213> Glycine max

<400> 530

gcagaattta gtaatgaccc actaacctag aattaatata acttaatgcc attaacctag 60
ggaattaaaa caaacttaat ggctgagtgt aactgaaatt gtggcaacca aaagtcaccc 120
ccaacagcca acaagtcagc caccatgg tctccaaaa ggctgatgcc taggttgcca 180
attggccct tattacaact cgaactaaag cccttttagt t 221

<210> 531
<211> 440
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 531

tctgaacatt agctagagcc aggtcaagag gtgagctcg aatccctgct agaaggtaact 60
gttctgatgc taanactatc tacttgctcc tttataataat atggaagaac tttagaatct 120
agttcaatga tcggcgccat tatattgaaa aacatgaatg ctctcttct ttatctctcc 180
cttgtatata atctctaata actaatttgc caatatgtat cttgccattt attgatctt 240
tagatttgca ttaaattggg taccacttgc gtgaaagggt atttgggtct tttgttttg 300
ttggtatctt attgctatct aattcgatgc tctcataaaag aactagtgc ccaattgcat 360
catgtctaac tgctattatt tatagccatt gcgaaacatg aatgctctct ttctttatct 420
ctcccccttgcata tataatctct 440

<210> 532
<211> 403
<212> DNA
<213> Glycine max

<400> 532

tagacggcaa tttcgagcgt ctccatatac tacggactc aatcagacat ccgagtaaaa 60
agttattgtc gcttgaattt gcctacaggt tctacattca atatcgagcg tcccgtata 120
ttacgtcact gaatcgac tccgagtaaa aagttattgt cgtttgaatt tgctctgagc 180
ttcaacattc aatttcgagc gtctcgatattt acggac tcaatcagac atccgagtaa 240
aaagttatttgc tggcttggaaat tggctcataa gttcaacattt caatttcgag cgtctcgata 300
tattacggga cttcaatcaga catcgacgaa aaaagttattt gtcgcttgaa ttggctaaag 360
gttcaacata taatttcgag cgtctcgata tatttcggga ctc 403

<210> 533
<211> 448
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 533

tattaaggga tcatttgcc tttggagtgt tatacgac tttgaactgc ctaaagtgc 60
caagaagtga gatcaaaaact aaatgcatga gcatgtctt gcccagctct agcttaagcg 120
tttaagttt gatgccaagc tagacatttc cattatgtac tcccttat tatttttc 180
cttatacttc ggagattcat gatcataaat gttattatct ctgccttctt gttttggca 240
aagtattgtt caatttcctc aagaaatttc tttacattt caccctcaca aatagagccc 300
cgaaacgctt atggaataga acacttcatg gtcataatgc acattctatt ggaacgatcc 360
aatttctcaa ttttggcctc attagaggtt ntcaagtgg atcgttcact tactaaaaat 420
agactttcaa catcggttat taatcgat 448

<210> 534
<211> 398
<212> DNA
<213> Glycine max

<400> 534

ctgatgcaac atttggagag gttaatgaaa caacgagatg atgcacttca tgagaggttg 60
gatcaaatgg agaatataga tcataatgga gaagaaagga ggagaagagg gaataatgg 120
gttcatagac aaaaccgaat tgatggtatt aaactcaaca ttccctccctt taaaggaaag 180
aatgatccgg aggccctactt gtagtggag atgaaaatag agcatgtttt ctcatgcaac 240
aactatgagg aggaccaaaa ggtgaagctt gtcgccccgg agtttccga ctatgcttt 300
gtgtggtaa acaagctaca aaaggagaga gcaagaaatg aagagccaat ggttgataca 360
tgggcggaga tgaaaaggat catgaggaag cggtatgt 398

<210> 535
<211> 405
<212> DNA
<213> Glycine max

<400> 535

tcgtatggtc tgaaaacaggt aaaagggcat ggtataagga aattgacagt tattttctaa 60
aaagaagggtt ttaagaagag tgaaaatgaa gtcactttat atgtgaagtg ataaaaaaat 120
gaagtgcac tcattgtttc cttatatgtt gatgatttat tttttatata taggaaatca 180
aattccttaa accaattcaa gaatagtggaa accttggaga atctttatga tagatacaat 240

taatgtataa gaaaatgatg gagagctacc atgtcgaaga tcaatacaat aaatataaac 300
aacatttgac agcttctaga aagagaaaac aatcaaaagg caattggtgt aagtgggta 360
caaaaacccaa gtaattctaa tggtcgcga acaactcact agatg 405

<210> 536
<211> 405
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 536

tcggctntca atgtcgagca tctcgatata ttacggact caatcaaaca tccgagtaaa 60
aatttattgt tgtcagaatt tgcaactgagc ttctgtttc aatttcgagt gtctcgatat 120
attacgagac tcaatcgac atctgagttt aaagctattt ctctttgtat ttgctacgag 180
cttccgattt caattacgag cgtctcgata cattatgggt ctcaatcgga catccgacta 240
aaaagttatt gtcgttagaa tttactcata gcctttattt taaattntca acgtgtcgat 300
atattacggg actcaatcggtt acatccgagt aaaaagttat tatcatattga atttgctcag 360
agcttctgtt ttcaatttgg agtgggtcgataaaatgtgg gactc 405

<210> 537
<211> 452
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 537

tgaagagcct cctcaatcaa actgaaaaac ctatattcct.caatgaagtt agcacccatc 60
ttttcataaaa acttgatggt gagaatattc caatcaagca taacccactt gaccctttg 120
cacccattt ttagggcttg ct当地ccacc acagagagca acattctccc aagcccccttc 180
gtcttataac actccctcaa gaacaagttc tccatgtaaa accctcgctt ctctagaacg 240
agagagaagt tcggaaaaaa caagacaaac ccaacaatgg aaacaccctt gagggtntt 300
aaattgttagt ttctcgagat aactattact taatggaaat aataaatgaa taattaataa 360
ttatggacta aattataatt gggtaaattt ggaagcagtt ttagagaaaa ctattatgg 420
attggagtag tnggtataag ggtcaataact ca 452

| | |
|-------|-------------|
| <210> | 538 |
| <211> | 390 |
| <212> | DNA |
| <213> | Glycine max |

<223> unsure at all n locations
<400> 538

ngaggagacg ctgaatcagt tgatgcagat atccatgtcc aactatacggagtc 60
ctccaacagg aacccttagaga tacaagtggg acaattagcc aaacaaatgg ctgaaagacc 120
cactggcaac tttagagcca acacagagaa gaatccgaag aaggaatgta gggcggtgtt 180
gacacgaagc caaatgagag tgcaaggaga agcagagaaaa gctgaaggag accagtctga 240
ggaaggaagg gcagacaaag aagaagagaa ggaggaagaa gagaagaatg tcttaatctc 300
tatgaccaan atccagctag cccaaaggagc tagaaagaag aaccaccagc cccttctaag 360
gaggcctncat atccttttagt actatcgaag 390

| | |
|-------|-------------|
| <210> | 539 |
| <211> | 334 |
| <212> | DNA |
| <213> | Glycine max |

<223> unsure at all n locations
<400> 539

| | | | | | | |
|------------|-------------|-------------|-------------|-------------|------------|-----|
| tgttcttgta | tgctctccctc | cacctagatc | cttaatttag | ccagtaaaaca | agtcaaagca | 60 |
| gtttggattg | caattgttga | agatctgatc | ttatggaaat | tgttgaagag | attgtctgag | 120 |
| gcggatgaat | tttcgatga | ccgtgatttt | cttgattaaat | ggcactgtaa | ggtctatgtc | 180 |
| ttaatagtga | cagctttatt | tttatnttatt | taaaaaaaatt | gtgcataat | gtctcattga | 240 |
| cataggactt | ttccttgttt | gcataaaaact | ccttacatt | tacgtagaac | tctctcgcat | 300 |
| ttgcataaaa | tgttagcata | tataactatt | atgt | | | 334 |

| | |
|-------|-------------|
| <210> | 540 |
| <211> | 321 |
| <212> | DNA |
| <213> | Glycine max |

<223> unsure at all n locations
<400> 540

caagcttaat ggtatggttgc cttctgttnt acggttcttt gaagggattt ctatttatcc 60
tagagaata ttggtaggatc aatcccgtcc ttgatgattt gcggctgttt tattcctctc 120
gtttcaggaa aaactaaact atntatatgg gtcttttct tttccctctt ttattgcattg 180
gtttgttaat ggtatgtacac caaagatagg gaaacaagtc taaagagagt cattgaaaag 240
aaatccctac ttccacggat gatctatctt tcgattaaaa gcgcttcagc atccatcaag 300
gagcatgttg aggtcaatgg t 321

<210> 541
<211> 213
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 541

attcattatg cgatataatg tgctgttaacc cattactaac caattcacat tattaagtac 60
tcgtctggta atcatgacac ttgttggtcc aacaaaaatc atttactggt gcaacataca 120
tgattgtcat aattgacaac acataatgac atgcacatgcgt attanagttt gagcgcgaca 180
cacattgact gacttgacta cacattctga gtg 213

<210> 542
<211> 394
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 542

caagcttga gccanaatcc tgactcacca tanaccttga cccagggtga gaatgccaat 60
ccctaccctc ggaagcaaaa aaagaataga gggaaattt ccaatcaaag aaaaagagaa 120
tgaaaatttc caatgaaagc aaaaaagaaaa tgaaggaaaa ttccctcaatc aaagagtggg 180
agaaagcaaa aaaaaggaaaa agaaggaaaa ttccccaaatc aaagagtggg agaaagcaaa 240
aagaacagaa aggaatattc ccaatcaaag aatgggagac agtataaaaaa ggaagaagaa 300
gaagggaaaga atgctcctga tcaaggatcg aaagaaacca gaagaaatgt gcagagaggt 360
cttggacca tacaatatct gaacagtaca gaat 394

<210> 543

<211> 500
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 543

agcttcttga accgagtagt accgtgctat gctactgtgc gcaacatcg accacttgct 60
tgtgcttgan cgactacaca ttgtcaagat gttgcctgag caagatgaaa gccttggaaag 120
aacgaagtgt gcctctgtcg ttgtggatga tatcttcaga gatacctgag tgaactgtgt 180
ctgaggatat cagacacacctt tggactttca aggagtttag tcttatacta tgnatatgca 240
tatacgttag aatgaagaca agcatgagtg accatggctg agagtgttag aacagaccgg 300
ttactgcatt gtgatcatgt gaaagcatga ctcatgagtg ttatgcatac aatacaccac 360
atcatagagg catatctgat atgaaacaca tgactatgca tgatgctgct acgggtatgc 420
ttcatgccaa acacattcgt atatctctcg gtgaagcatg acacagcgtg gtcatcgaca 480
cagagctcac tgtgagaggg 500

<210> 544
<211> 375
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 544

atgaagtaac cagctcgccct gagcgagcaa gttactttcg gatgaagcaa gcagctcgcc 60
tgggcgagct actgtgcaac ctctaccctt catttcctat aaataggcat gaggggggct 120
gaaggaacgg tccaacattt gaaatcaaga ggatttagaga gaaatttgcg agaagatgga 180
aaaaaaaaga agaaagataa aggttgagac gctttcgtaa cgtttctgtg atcgattccg 240
agatcatttt tcacgttct tcgacggat agtttctatt attgaagcta tgaattcatt 300
ctatgcaccc ttagggacc atacttgctt tacatatctt catcttcatt ctcttaccat 360
tagngatctt tcttt 375

<210> 545
<211> 276
<212> DNA
<213> Glycine max

| | | |
|---|---------------------------|-----|
| <223> | unsure at all n locations | |
| <400> | 545 | |
| acaacatcct anatagatta tattatattg tagcananat taanaatann tttaaattta | | 60 |
| agagagctng ataataataaa aataaataag aaaaagaaaa aatagaatc ttactaatac | | 120 |
| taacagtctc tcaataaaata ttctacaaaa attcttgaag atcttcctaa acatttgaa | | 180 |
| agtcttgcaa aaacacaatt ttntntaaaa ataaagtaca nnatttgaa nacaagatca | | 240 |
| ctaagaattt ggaacttctt aagtccaaag atcgac | | 276 |
| <210> | 546 | |
| <211> | 428 | |
| <212> | DNA | |
| <213> | Glycine max | |
| <223> | unsure at all n locations | |
| <400> | 546 | |
| atgcaagctt gtatattaga caagtctaa gtcatcatgt tcatanaaca aatcatttg | | 60 |
| ctaagtcaact ggcatctaga agtcctaatt ctctcgtaat ggtgtagaac gaatctttgt | | 120 |
| gtagtggttc tgtgaagata tttgcgagtt gantttgggt atctacaaat tctagaacac | | 180 |
| agtcacctt taggacatga tctctaagaa gatgatgcct aatttctatg tgctcggttc | | 240 |
| tagagtggaa aactagggttc ttagatatat ttatggcgct tgtgttgtca cattntatgg | | 300 |
| ngatgtggtc taatacataatc ctataatcag atagttgtta tttcatccac agaatctgtg | | 360 |
| cacaacaact accagcagag atgtattctg cttcggttgt ggataatgct acaataattt | | 420 |
| tgcttctt | | 428 |
| <210> | 547 | |
| <211> | 298 | |
| <212> | DNA | |
| <213> | Glycine max | |
| <223> | unsure at all n locations | |
| <400> | 547 | |
| acccagtcac cctcattcag aagtagctct ntcttcctc tattgcctnt agttgaatac | | 60 |
| acctttgttt ggttctctta ttgggtctta accctctcat gcaacttctt tacaaaactct | | 120 |
| gaccttagatt ccccttcttt atgtataaaa gaagtgacta gtgtgaggtg aatgaggtct | | 180 |
| aacgggtgtta gggatngaa cccatagaca acctcaaaag gggactgctt ggtggttctta | | 240 |

tgaaccaccc tggtaggc aaattctaca tgaggaagat actcatccca agacttat 298

<210> 548
<211> 376
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 548

ttcaacaagt gtatgaaatg catgtacttc tttatgtga gaaaccactc ttctcgctg 60
acatgttcaa cactntagct ggaaaacact tctttgtgag acagagcaag tctatgcaac 120
aaagttcttc ttttgcgtt gattgaggaa tattagagct tggcttcatt tattcttcatt 180
aagacttggc agatcctact cgaatgtctc tacaaaatag atgtagaca caggattaaa 240
tgaagtctta aatgtcaact ttaatatcga atcagatcat gattccatct tgcaatcg 300
cgaaaatataca gacgttagact ctgcgaaaca tgcgttgata acgcataatga tgcaatcctc 360
ctaaagatgg acccat 376

<210> 549
<211> 273
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 549

ctgatcaaca catgcacagt ggccaaggat gcatggaga atcctgaaac cactcatgaa 60
ggAACCTCCA aagtgaagat gtccagaatg caactattgg ctacaaaatt cgaanatctg 120
aagatgaagg aggaagagtg tattcatgac ttccacatga acattttga aattgcaat 180
gcttgactg ctttgata aaggatgaca gactgaaagc tggtgagaaa gatcctcaga 240
tctttgccta agagaattga catganagtc act 273

<210> 550
<211> 339
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 550

gactaccaag ggacatcgga agaagatcgc tcatgtgccca ttatttctt cctatttctt 60
aacccctttt gcaccattta agtattgatt aatcttaatt gtcaaattaa ttaagcaatt 120
ntattatttg gcccattca gctcaatttg atgtttaat ctaatttcac gaattaatga 180
agcattgcgc ttgaatccag aaatggcctt ggacttgaag agggcagact attntattct 240
acaaaattnt atcttatcta gactntatct tatcttagata tttatttagaa ttgatctcat 300
ctagatacta tttcatctag atcttatctt atcttatct 339

<210> 551
<211> 322
<212> DNA
<213> Glycine max

<400> 551

agtgcacatg ggcctgtgtc ctgcactaa gcctacgatg ctgccttagc acaagtgcct 60
gtattcgcgc tgaacgcggc ttgagatgtg ctttcctcgc gcttagcgtg tgcttctcga 120
tgagcggct ggcactgag cagacagttc tcactaatcc tcatgtaaaa ctttacctt 180
tatattggtc tatactacg tcttttatt tgatccctc ttttatatct gcgatcatag 240
aaagagaact gtattttaaa ataacataat aatgctaaaa atactttaag gtatgttata 300
taagaaaact atattacatt at 322

<210> 552
<211> 286
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 552

ggagatgaag atgattgana agtactgtat ggattgtgtt tntcatggcg aatatgtgaa 60
tgtatgtata catgattntg atgatgtcaa aagaagaatc aaacaaggct catttggattc 120
aagattaata caagattgtt tcaacaaata aagccttgat tcaagattc ttcaagatca 180
agccttgct canaatgaaa ggtttcaagt catccaaggc acatgtatc gattaccatt 240
acatgtatcatt gattaccaag gcacatgtaa tcattacaa tacatg 286

<210> 553
<211> 369

| | | | |
|-------|---------------------------|---|-----|
| <212> | DNA | | |
| <213> | Glycine max | | |
| <223> | unsure at all n locations | | |
| <400> | 553 | | |
| | | agcttgttcg gntgatttct ttgataatgt tttgtcttt tgcttagaag gatttcttat | 60 |
| | | tntccttga ctttataatg gttctttgtt accgactaaa tttcctcatt tatgagggaa | 120 |
| | | aatgcgtct gttttatgtt ctggtagat gagttcgtgc caagcttgaa tgaattctta | 180 |
| | | agagggtgtg ttttcaaagn ttatacttta ctctgtatcc tcttggtaat ttcagggttc | 240 |
| | | aagtgccatc ttcaagcgctg atcttttgg tgactcaaga gancacaaca tcgatcttac | 300 |
| | | tgctggngac cttatcaatc gattatcttt ccaggtattc tttacctttt actcgaattt | 360 |
| | | ggaaattca | 369 |
| | | | |
| <210> | 554 | | |
| <211> | 526 | | |
| <212> | DNA | | |
| <213> | Glycine max | | |
| <223> | unsure at all n locations | | |
| <400> | 554 | | |
| | | aggggnnnnn nnannnccgc tgtcatgtnt tccatttatt tttcanctcg gnancancca | 60 |
| | | agacnnaagc nanatcctca gatggctcnt gtaggactag acttataccca acctacacca | 120 |
| | | tgggcacaac atgttgcaat cgaaaaactga gtccgcacac ccctaactta agactaagt | 180 |
| | | gcagtgtatgc ttcatgcaag tgctggggca acagtagtatt taccaatgct aaagtgacgt | 240 |
| | | aactaggcac acaaatggat gatcataccg agagcataca aacattaaga actgaattaa | 300 |
| | | gcattgaaca cagggaaacac agtcaactag atgtacaagt aatgacatta gactatctac | 360 |
| | | agaaatcccc agcaagggtg ttcaagccagc cattacagaa aagctctaac agtgatgaga | 420 |
| | | ttacaaaacc taggcctntt tgcgaaagat gctccacttg ctgcctctag agcggtattt | 480 |
| | | cgagataaga gtagggcgcg ctcttgaatc attgcaaagc atctcn | 526 |
| | | | |
| <210> | 555 | | |
| <211> | 320 | | |
| <212> | DNA | | |
| <213> | Glycine max | | |
| <400> | 555 | | |

agcttttact cactgtttc atcagatatac atttcccttc acgagataat cgccccttt 60
cagatttctt ataatgtcgc aagaatcagc agctccgaat tctgtatttc gattctactc 120
ttgaagttagt attgacacat gcccttcgt gacaattta ctgctatcgc cattcttatg 180
ctcccataat atgaggctat actatgctta tcttaactcc gaacagtgtt gcgttatatt 240
tatacaaata tataccattt atactcacag tacgtatgtg aggggttaat tcctctaact 300
cataaggacc attcgaatag 320

<210> 556
<211> 330
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 556

gctctagccc tcgaattgtc aacataactt cgctagagaa actctagcga gtacatcgtg 60
catatggaga acatctngta nagcccctgc acatagacca aataaccatc accaaactat 120
ataattatgg tggaagtgtg gaacaacata aagatggcaa gttattaaaa tccataaagc 180
cttataatctt agcagaaaaa ctcagaaatt agttaagcaa acatgttcca catatatttt 240
gattctatgg tttattaaat aacagaagtg gagtnctct tcatgcacag tacacacttc 300
aattatgaga ca~~g~~atgttaa gtaatttaag 330

<210> 557
<211> 303
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 557

agctntgttc ccttgcttg gatgttgaa attctccaaa ttatgtactt aaaaaatgtt 60
catacggcgg gttttggagg ttcacgtgat ataccctttt cttattccca aatgagagga 120
ggccttactt aaaaccttcc cagttccct tccattgcta tctcatccat ccaaacatat 180
ctagctcaat gagaaggat ccaggcttc attaactatc tagctgttat acaattgtt 240
tatgttcaa tgtttttgt gtatttctt ccattntctt gcccnaact ccacatgtt 300
ctc 303

<210> 558
<211> 213
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 558

gcttattatt attccaataa ctntgtgtgc ggaacaataa aaataacact cataaacata 60
ttaaaaagca tttaacaatg aggaanaaaa tgtcatatac caaacaagaa gaatagccac 120
aagagaataa caataaaaata gtatttattt tctaatcta cctncttatt acctaattag 180
ctcaatctcg caaaattgaa aatgcacaat tga 213

<210> 559
<211> 311
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 559

tgcgctaaggc cactgctgtc ttgcttagtg ggatggtctc actaatcgca tggttcaggc 60
tttagcgagac aacactntnt gaaccttcat aatntctcc ttttacttg aanatgaagt 120
gaaatttaca ttaaattgaa taggaaggct tctagtgagc acaaattgata actaaactag 180
aaatatttac aatcctacca aaaaataacc ataaattggg agaattatnt acattntgga 240
cactnttcta tacaaaaatt agtcgtaaaa gacgactaac acatagtcat atatgttggt 300
atgttaagtag t 311

<210> 560
<211> 375
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 560

ttcttaataa actnttaatt gggtgcaata tttatgtga taaatttattt cttttgaatg 60
atatattatt tgataaactt aatanaaaaat ttatataatt taagaaataa attgtataat 120
taacataaac atatttgcata aataaatata tcataataac tctcaatatt tgtcttacga 180

gaaatataaa catacattca ctttctcttt catctcattg tgcaaacatc tctctattta 240
tttttcatta gactacttat actctattac ttatthaata ttgagaatta atgtgcgaat 300
aactcatgag aattctatat aaatatactt tatactatga aaatatattt caaatactta 360
ccggcttggaa aatga 375

<210> 561
<211> 335
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 561

tgtatggaga tctgaggact tcttctctct atgtcttcac aatcatctct taggcggaag 60
ttgcagaatc atcatgatga agatactata ttcnngcacgt taticgttgat atttggacc 120
cttaccttga ttgcattgct taaatatgta ttcatactat tgtgtgctga tgacaacgg 180
gaacgtatgc tgctgataaac tttattctct tagataattt ttcttataaca taggttacga 240
tctgtgtgag tgtgagctga gtatgatact tgtgtcatca ggtggAACAT ttgcctttat 300
tcgctgctgtt ggatgcatgc cacgcttaaa ttact 335

<210> 562
<211> 298
<212> DNA
<213> Glycine max

<400> 562

gatctaagaa tagaaaaact taattatcct acttggatga atatgaagct tgaggaacat 60
ggatagaata agaatgaagg aagaacccgt gctattgact gtttgcctta catggacaaa 120
tttatcgccct acttaactat gtcaacactc aaccaatatt gattcttctc attgccacc 180
accctaccag tcaagaacac ccaatcatcc acaaggggcca cccctaaatc agccgcacag 240
ccctgctgtc ggacatacga tatcaaacac cactcttaac acataccatt acactaac 298

<210> 563
<211> 270
<212> DNA
<213> Glycine max

<223> unsure at all n locations

<400> 563

cttcattcaa catgtcatat gaacaggana gantatacaa attcattaac aatcaagtca 60
cactaaacat tacaggacaa cataagccaa cctaaaatcc tagaatgcaa acctaaaaac 120
cagtctctga attgagcaga cctaaaccct aaacatctaa cttccaactc tggaaagccc 180
agaacaaact tcccaaagat caaatccaa acccaacctc agaaccaga aacgtatatac 240
tagctacatc atagacaaac agatgacagc 270

*

<210> 564

<211> 339

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 564

tcaagttca gactttcgat tcggagtaag ttccatttct tgagattgtat tttattatac 60
ganagtaaag gtgagggtca ccaccgatat ttatgtttga attggataaa gtcattaact 120
ggtgttattgg actaaaatac tattaatata ttttgacata acaagttgtac cacatgctac 180
aagattatgg cagaaattat cctgatagat ttttaaaga tttatgtatag atttatagtt 240
cattatattt aatggattat atgtgagaat caaaacattc gaagtattcc tatgaatctg 300
cacgatctga aagagttta ttaattctct ttaacaaa 339

<210> 565

<211> 515

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 565

aggggnccat gacccatatg atangcttag natnttcaa ancattttgc aagaaantct 60
gctangacta tataatgca tttttgtgtat tgaaaacaag ttatcagatg aagcttaggag 120
taggtggagg ggggtttaac tagtaattac cgcttttatt cacaattaat tagccttata 180
tatttttcg ggggacgggtt ggtgctgggtt cgaatcgaat gaaatctctc ttctgttagtt 240
tgccaaaaat agtttcttga gatcaattttt atatttttta tttgtatattt cattcttgg 300
gaaccgtgaa ctacatcacc atcttctatg aataaattac ctctaattta tgggcttatt 360

ttctcttgt ctataaaaga ttaattatga ttcttaataa aataaatatt tttcttaaag 420
aaataaccac tataacatta ttagatcctt taagaatgtg aagagacttt ataacttagg 480
cgtatcttag agtgcattat ggaatttatac caaag 515

<210> 566
<211> 206
<212> DNA
<213> Glycine max

<400> 566

gctagacatg aggaagcggt caagggtgaa acttcctgct tttattgttg accacagagt 60
ggtagcttggaa gatatgtcgc ggaggtcacg agaccttgtc gacgtcaggt ggtgtgctat 120
tgcccaaaac caagcttgac cacatcccga cccaacccgg gcatagtggg acagtgagaa 180
cctgtgatgt acctaaggcag gcgagc 206

<210> 567
<211> 314
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 567

ggttgaatta agatatcaca tactntntct tnaataanaa atcttatttg attntaacc 60
anatcccaag aattctttca naatgaactc ctaaataatt atgcannata aacttactga 120
atagaagcaa taagcaataa ccaataaaag agttaaggg aataaagaat gcanactcag 180
aattatactg gttcgccccac atccttgc ctaagtccag tcccaagca acccgcttga 240
gagtncacta tcttgcnaaa gcccttacag tctgaacaca caggacaccc ttcttggtc 300
agattgttac acaa 314

<210> 568
<211> 340
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 568

tgtgtcgct gcatactgca atatgctgac tntgacttca tttctgccca cattgaaacc 60

ttgaaacagg ttnttatcca ttgcttcttt cnaccagcgc gttggtcctt ccacaagaat 120
gttaaacagg aacggngata gagggctcc ttgtcttaag cctctntggn gaataaaactc 180
aggtagga ctccgttaa ccaatatgga tatggaagca gactttatgc agccttcaat 240
ccaagttacc cacctgtcac anaaaccat cctcttcaac aaatata tagta caaactccc 300
agataactgaa tcatangcca ttntataatc tactttgata 340

<210> 569
<211> 240
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 569

caatttgaag tgatcatgga ggtgaatntc ataatgagtc ttttgaacac ttttgtgaag 60
aacatggaat tcaccacaat tcttattccc caagaacacc tcaacagaat ggtgttgtgg 120
aaaggaaaaa tatatcccta taagaaggtg caagaaccct tctaaatgaa acaaggttac 180
cgaagtactt ttgggcagat gttgtacata ctatntatta caccttgaac agagtactta 240

<210> 570
<211> 351
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 570

tgeacaagac tcttaatat tgaagagtat cttgtggaa ctttcttccg acgaagacac 60
tgacaaaaac ttatcttctc cttttggac aaggtatgtg caggctgggg caagtaaatt 120
ttcttccat cagaccttgg atgcaattgt gatcgatac ccatacagc tagatcttga 180
tgggtattca agccatcctt cgtctgcct tgaatgttaa ggagcgtncc aatcacattg 240
tcacaaacat ntcttcac atgcataaca tcaatacaat gtctaacgtc aagatcacac 300
cagtacagaa gatcaaagaa natagacctc ttcttcataat gcaactctga c 351

<210> 571
<211> 421
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 571

agctntgaat tctagtaaaa aaaaactcct catacatatt ctaatactca tgcacatttt 60
acattcaaaaa ctggaaactt agattcctag ccatgagtca tcctttggc actgttagtt 120
agcttctaca aactaccac acactcacaa tgtgcacaat ttgttcgca agctaaattc 180
cacaaaatca tccgcaaatg ccattgaggc atttcaccga acacttggtg ggcatatgtt 240
taagcatgaa aatcaaggga atgaggcaa tgtggcttgc cccattatct cagaatgcac 300
cctatgccta aggccatacn ctacaacccc acaattcaac aaaaacaagg aaattcaagg 360
atacateccct tcacgtttga gcaaataatat gcaacttaga gcaccaaaaat atatcaatgg 420
a 421

<210> 572
<211> 232
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 572

atatcagtat aactctgata tggcttacga ccggactcag ctgcagaaca tggttaaaaa 60
ggatagcgag tcctttaag agtacgctca gctgtggagg gacctggcag cgcacgtagc 120
ccctcccatg gtcgaaaggg aaatgattac catgatggta gacaccttgc cagtggttta 180
ctatgagaaa ttagtangct acatgccctt cagttcgca gacttggtat tc 232

<210> 573
<211> 317
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 573

atctagaggt ataccttagga tctctatcag acactatgct agatggcaca ccatgtaata 60
tgacaatctc actaatatac agacaggta actnttccaa ggaagatatg atattaatgg 120
gaataaaagtg agcagacttg gtcagcctgt caacaataac ccacatagaa tcaaaacctc 180
ttggggttct aggttagtcct acgacaaaat ccatagaaat atatgtccat tgtcactggg 240
tatcttcaag gggtgtaact atcctgaagg gctctgatat cttatactta tgacagacta 300

aacatgcata cacaaac

317

<210> 574
<211> 180
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 574

agcttgaaaa ataaatattt aataaaaata tatttattta ataaaaatata taattttgt 60

ccaaaaaaaaat tattaaacta aaatagggtgt taatttaaaa ntgggcttc tgccctaaat 120

aagctggacc ggccctggaa tgaataatg ggaataaaag aaatttggtg aaaagtattc 180

<210> 575
<211> 380
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 575

gctgccactc aagagacatc tcacctatat ctnntattaa aactatcatg tacatctgtc 60

cattattcan aaataaacacc atctaagcaa acttaactga gtagagacta gtactctcct 120

tcttccatac caatatgtcc tcctcaatca gaatcaaatt aatacttcta aagtcatgac 180

cttntatcta agtaaattat tatatntatt ttcctaatg atatagagat ttctncattc 240

ccatcaccac aatcaatntc cttccctccg gtcacctaataa ctttgacacc ctggctttat 300

ggctactaca anggtgtata atcttagtata cttcctatat gtggaaataa aaccacttca 360

cccgaacgtc ttgagaaaaat 380

<210> 576
<211> 147
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 576

tcaagcttct tctggacctt gaacaagcaa tcaactttct ctgtcacaaac catgctatgt 60

gctcgcgact ggtcccttcc ttcccttcgc aacttgagct cactattgct accccataga 120

gctncgagaa atttggccg gccatac

147

<210> 577
<211> 324
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 577

caccggcac gtgtctacta tcattgttat aatctctntc tctgggttgg gggtgctact 60
ttagttgcca agcctctcca tctttngcg tggctttga aagatccgtc cnccctattg 120
cacacgttct gtagttgcat cctatccgaa gacattatac tgacactgcc taacgaaggc 180
caccactagg tccttccaag aatggactcg ggaaggttcc aagttagtgt accaggtaac 240
agctacccag taagacttcc tcaggagaaa tgtatcagca gtttctcatc ttttgcgtat 300
gcacgcacatct tccgacagta catc 324

<210> 578
<211> 215
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 578

gatccanaat cctgactcac catanactct tgacctagg tgagaatgtc aatccttatac 60
ctcggaaagca aaaaagaaaa gaaggaaatt tccaatcaa gagaaagcag agaaaaaaaaa 120
aaaagagaag ataggatatt cccaatcaa gagtgggaga aagaaaataa aaggagacga 180
aagaattttc ccaatcaaag aatgcgagaa agtat 215

<210> 579
<211> 270
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 579

ccaaaactca ttccgtagat ccctcttgc agactaagtt tcaatcttgc ttcaatcaaag 60
ttctaaggca acagtacatt tcccaatgct aaagtcaccc aactatgcac acaaatggat 120
gattagacca aaagcataca aacattaagc attgaacaca aaaaacataa tgaatttagat 180

attaagtatt tacatcagtt gctcatttga aatccncaac tagggtgtnt agccagccat 240
tacagaagag accctaacaa taataagctt 270

<210> 580
<211> 440
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 580

gatcaagct tgcctctaca gaggcatata tacgaanaat aatccacaca cacttgtaca 60
aataaaaaga ataataaatac cacagcaaca accttatctc tgtagccgtc aacaccaatg 120
ggcgaggtct gtataaccat tctctttcc ttttctttc ttcaattacc atcaatgtat 180
cattccgggt tctgattttt ttttgtgtta taaatacgaa gagaaaaaac tagagggaaa 240
caaagtggaa gagaaaaaaag cacaggaaga agaaaaagaaa gaagaagaga caaaacagag 300
gaatcaaaag atggcaagaa atccaacgag gaatctgcgc caccagaaat cgtgcaaggc 360
acaacccctca caatgcata acactttaag cacgattct gacatctttt ttaagntaag 420
gactaaattt gcacacttat 440

<210> 581
<211> 368
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 581

gagggagaga cagagagagg ttgttagcag atatttaatg aagannaagg gagagaagtt 60
gaactttgag ttgtgtctca caagactctc attcatcaga gttacaacaa gtgttacata 120
tgcttctata tataactat gtagcttcct tgagaagctc tcttgagaaa acttccttga 180
gaagcttcct tgagaaaact tccttgagaa gctagagctt agctacacac acccctctca 240
taactaagct caccccttg agaagcttc ttaagaagat tccttagagaa gctagagctt 300
aactacacat accttctaa tagctaagct caccccttg agatgagacg cttagaactta 360
gctacaca 368

<210> 582
<211> 307
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 582

gtcgcaacgt gcccttcgcf ggcgaggcgat ggcgaggctc acgggtgcgc tttccaaagg 60
aggaaagatg cgccggagtgc ccaccaacgt ttatttgtgg aaaacgtctg ataaaaccga 120
aggaaactgg tcaaaaggaa aattctaagt tcgggagttg tatttacgct tgaggaaggt 180
attagcacct cacacgtttg tcccatacgaa caacagtcta ttttttagaa ttgcggaatt 240
gtgttatctt aaccttttagt tcttttatac ttttgaggc aacanaagcg gggctttgc 300
tcctaca 307

<210> 583
<211> 361
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 583

tgcggtaaa tgtcgccaag gtagatataa tttagaagccc tcgctggtga attagtgaac 60
ccagaatcat aagagaatat gaagatggag gaagccgcag gcagtggcaa cagtgacggt 120
ggcaacaaat ggtggtgng agtggccagc gctgctcana tggaatgng aattcgtacc 180
ttcgccaaag ggcacggagg cgattcgcgc ctcatgcctt tcanagcctt cgtttagct 240
tccctcttcg tcagcagcgc cgccttcgccc ttctttctac tcctccaagc taacggcattc 300
cacagggtag tactgctctc atcttcattc ccatttctgg tttctaactc tctntatgtc 360
t 361

<210> 584
<211> 201
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 584

catcatcatt gtaactatag ctatccatta taaaatagct tagacaatct aggtctgctg 60

cttctaaacc ttntatttat ccttgaagca gctcaatgtt ttttatttcg cgtcaactag 120
tgaagagtga acactgaagc tgttctgcag actcattgat ttccttcat tgcgacagaa 180
cctaaccata ccctgcccgt a 201

<210> 585
<211> 198
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 585

acacacccct ttatactaaa tgcancccc tttctatTT tgTTTgaaat tcTTTntccg 60
taacgttacg aaactttacg aatctcgtaa cgatacttat tntccttccg cacggttacg 120
aatccttacg gattatgtat ttactctnnt ttggcttca aagaagttac ggacactcac 180
ggattgcgca aaaacacc 198

<210> 586
<211> 399
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 586

agcttggtn tggtaatAG cacccacact gactccnca aggtctactg atcctcgcaa 60
catatctcca ggtaccactc tgggtcaac aaataatagt aggaagattg actcttccat 120
gcttctcac atcaagctta ttggattatg ggcacccgt catatgttgt actagggtggc 180
aatcgggcga tggcacaaat caactatcac attccacaa gccaggcata agcacaccat 240
ccncagctgc ccaccttAA attagctca cgtgcattGA cgtagccttc tcctcgTTcc 300
tctcagcacc gggccccat caacccctc aagctntcac aatatccaaat caattcaatc 360
ccatttgta tggaaactacc ttAAacAAAG aanaacaga 399

<210> 587
<211> 314
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 587

gcttgaatcg atacacacat actgtatatcg attaccagag cagtatttca gattatattc 60
tcatcagtca cannttttt attgggttctt gaatggccat canaggctta tataatatgtg 120
acttgagaca ncgaattgct aagagattnt cagaacanna aggtcttatac ctcttanaaa 180
gcanaatcg ttcatcctct taanaattcc ttggccaaaaa cacttgtat tcaataagga 240
attagttgag tgctcaaatt gttcaatcta tctctt~~t~~caa gagagaatac ttcttctctt 300
cttcntatt ctga 314

| | |
|-------|-------------|
| <210> | 588 |
| <211> | 383 |
| <212> | DNA |
| <213> | Glycine max |

<400> 588

tgagagcgcg atcttatact gtgagagaac gactagctgc gagtaataat cttgcatca 60
atctctgaat tctagaatga aatgtataaa tgaggacatg atgaatgcta tgattgcaca 120
tacacaaggc ttttgaccaa aaagcttacc ttgaatgata attatatcct tcgcaccctt 180
tatgagctga atgatattgt caaagatttg aaccctgAAC ttAAATAATA acctccagat 240
accttggtag attctaggag agcatatggc tcaaggcaaa ttaccgcaaa tttgcggagt 300
ggaactaatg ggatgcaaga aagaataaaa catcgacaca acaacacata ttttgttat 360
aacaataaaag gagaatgaaa agt 383

<210> 589
<211> 425
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 589

gctctttag tggtgcttct tgcaagcnnc cttgttactt gtattgattg ttattnccct 60
gtgttgtcaa agccattcaa tttaacatata agttgagaaa ttgatcatca aatgtatgtg 120
catganaaaat atatatgttg agagatattg gcctttcat gattagttnt ttaatccctc 180
ccagaagcta ccatggccac ccataatana tgccattcta aatttttag tttcttacaa 240
attaagtatg gacaatggac ctaagtgaat ggatttgacc tacgatagtt gatataattct 300

tgatagttaa tatctgaaca ctcacttatt ntatacactn catcaataat catttttct 360
ctttctatca tatcatctat ctatantan tttttcttt tgggtcattc tctctggctc 420
tatct 425

<210> 590
<211> 318
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 590

agctttagg attatggtgt acccatcaca tttgtacta ggtggggc gggcgatgga 60
gcacaaaaca tttccacat ccacaatgcg cgcataaacc caccatccnc tttgcccac 120
ctccaactga gtcacgtac tcccacgtag cccatatcct cgtttctctc aacaccgggt 180
ccccatcaat cttcccaagc ttccacaaca tccaagcaaa acaacattcc aacagcacaa 240
gctatcacag ccaagcataa cagagcagag gcagaaaact ctgctcaaca catcaaccaa 300
aattacagct tttctcac 318

<210> 591
<211> 357
<212> DNA
<213> Glycine max

<400> 591

catgcgt tacattccct ttagcattca ttattgaatc atttcagcct ttgcttcgt 60
gtagcttagg aaaaacgc ca tgtattctcc tttctttctt ccaaagccat ttctaacatg 120
ccaagcactt tctccatcac ccacatccac cattagccac cacaaccat cgttgctctc 180
cggtgaaacc ccacaccgat aggaaccctt caaccaaagt ggaatcttac aacttggctt 240
gcgggttgg tagagaacga aaccctaatac tgacctttt ttttatcga gaggattgog 300
ttgaattgat gagcaacgac gaataagaat cttcaagtga cgcgacgagg aaccgc 357

<210> 592
<211> 172
<212> DNA
<213> Glycine max

<400> 592

acttctgctt tagtttagacc acatatgtac ctgagaatgt cgccgggtca ggagacttgg 60
gacgcaggcg ggtgctattt ccaaaccaag cttgaccaat cccgccccaaac ccgggcatacg 120
cggcagttag aaccttgatg tcctaaacgg cgactcctgg cagcaacaga aa 172

<210> 593
<211> 285
<212> DNA
<213> Glycine max

<400> 593

tcttcttgct tatggcgctt ctatggaggc tggatctttg agttcaatg gagtcctta 60
atggtgattt ttccacatgg agatgcagcg gaagacatag gataagaggt gataggacgc 120
accatctatt aaggaataag ccatggaaga aggagcttca ccaccaagat gatccttgg 180
ttacaagctt ggagattgtat gcttacttggaa ggaaaagata gagagaacga gggagcacga 240
catttgttga atataagagg gagagatgtg caactttgaa gtctg 285

<210> 594
<211> 495
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 594

tgcangtcaa cccctttaaa accgtggtaa ntncntttg gantcacgac acactatcta 60
gtactcaagc ttaatccaca nagtccacga ttttatctgc atctcctta aatgaatgcg 120
aagagaatga agttgtcat tttgcttgc caacaggcaa cccaaaggcca tgagaccatt 180
cctatcaaac aatcactcat tcactcacat acaccacaaa cctaattgg agtctcgaat 240
gcttatcctt tgagcactca cagtagatg cccctcctcc tactcccgcg agagtaacga 300
gagcatgttt ggtgaccctg tccaaagaact tacttcttct tagagacaca cagtcgtcgc 360
ctcccttcct gcaccgagag tacaaatgca gggtggtgag cttaccaaaa actactctcc 420
cggggccggc ctgggtggtc gacctgtact cggagtatgc atccgtgcct cctctgccga 480
gggatggacc tgccg 495

<210> 595

<211> 390
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 595

agcttaacca agaaaaaggc taacaatgtt nttagcaca attgaaggaa ataaaattca 60
gaattttagga attcaagtaa caatccttca tacaaccaat atattacctt aaagagattt 120
tttttanag ttcttcaagc atcaaccatt caacccaaat ttgtctctt ttntttttt 180
tttaattnt gcttatacga atttctgttt ttttttata acaaagagat caaaaggctt 240
aactttgca atggttcagc ctaaaaaaaaaa aacatgaaca agaaggtaat ataaatggca 300
aagaaaataa agaaggatgt tacccaatat ttccagcaaa ggaagtgttg atcctagaat 360
cggaactctg ataaccaaat gatatgaacc 390

<210> 596
<211> 318
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 596

cacaagttag tttattcgat tattagagtt atctcttat cttaggagag tgattctctn 60
attcttggtg atcaagacac tctgctgate naggacttca caccttgtgt gtgccttcct 120
gg nagagtgt tcttcttct atcatctcac cttgtctttt accacaattc agaaatcacc 180
ttgccaaata tctgtgacat actccattac acactcaata agtatttga cctaattgaat 240
tcaaacagac tttactcggtt gaataaccta tggacctgac ttcgtatgca tttatatttg 300
tcgcacacta ccacgtta 318

<210> 597
<211> 396
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 597

aagctctact aaataaggct agacattaaa attttgatag anaaagtac atgcatttc 60
cctgtatata gacaagattt gagttgagat gaactgaaga aagagagggc atattttagta 120

ctcttagctag tcaaagtggt atatataatgt ctaataacga acctggtgct ccataatttt 180
tgcacatcccggt gtcctgtttt gttgaatctc tccctccaac ctaaaagttat atagtcagtc 240
acaaaaaagag tagtaaacagg gttaagcata tatatgtgtg ttatgtcagt tatcttcac 300
aaagcagttat atgctatata tatkctcac agatcaaagc tgacgatagc taaaacttact 360
ccaataggc tcatgcaga acataactcat tcatgc 396

<210> 598
<211> 511
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 598

cggaaatgaac catgtagnac ccatcaactat tacgtataaa gtctacagaa ataccagtca 60
tgcataatctc caatcaatcc acgggcttat ataacatgca aggtgtttgg gccactaaca 120
tgaaccacat tcataaccca cctgtcaaga ttctgattaa tcaatgctgc tgcaaatctg 180
caaaaactcaa tatattgaag attgaagatt ctcataggaa aattcctaaa tgcataatcta 240
taacaagaaa gtaaataaga caaaaacaat accctccaaa acctgctcgc atgtccataaa 300
catttcttaa tctgatttct ttccagtgtt aaacacgaac atagcttgct attatttcat 360
tccagtattt cgaatctgcc ctaaaaagct ctgatctgga tgtgaaagca tcaagctnta 420
tgcttgaag cctatcatgt ggggtttgca aacgtgcattg ccattagtaa catttgctca 480
tatccgttct tatgcattca gagatgcattg t 511

<210> 599
<211> 415
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 599

gagggactca tggtcactat gaatgacaaa ttccgtggaa taaaggttgt ggtgccatgt 60
tcccacagcc cgtactaagg catacaactc cttatnataa gttgaatagt taagggtacg 120
accacttaac ttttcactaa aataagcaat tggatggcct tcttgcattca acacagcccc 180
aatcccaaca tttgaagcat cacactcaat ttcaaaagat tttgaaagt ttggcaacgc 240

aagtatggng gcattacgta gctttgctt aagaacattg aaagcttctt cttgtttctc 300
tccccatttg aaaccaacat tcttcttgag cacttcattg agaggtgctg ccaatgtgct 360
aaaatccctc acaaatcgta tatanaaaact tgctaagcca tgaaaactcc tcacc 415

<210> 600
<211> 451
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 600

tatttagcc gatgctctgt ctcggcgtca tgcattactt tctatgcttg aaacannaat 60
gattggctt gaatgtttga aaagcatgta tgataatgat gaaacttttg gagaaattct 120
taaaaaattgt gaagaatttt cagacaatgg tttctttaga catgaacgct ttctttcac 180
agaaaaacaaa ttgtgtgtgc ctaaatgttc tactagaaat ttgcttgatc gtgaaggcaca 240
tgangaggtt taatggtgca ttntgtggtc caaaagactc tatagacatt acangaacat 300
ttnntattgc ctcatatgaa aaaggatgtg cagacactct gtgaacatcg cattgtattg 360
taaaatgcaa gtctaatttgt aagcctcatg gattgatact ccattgcaat accgagtatc 420
ttgattgtta tcatggattt gtttggctg c 451

<210> 601
<211> 322
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 601

tgcgcatact tcttcacgaa cgttcacttg cacaagacat tcttataact atgtaaaatg 60
aacccatata ctattaatgc accttcgtta cctagattat ttacatgtac tatcaacgtg 120
tatgtgttac ctacatcaca cacatttct ttgcttagact cacatacatg catactctaa 180
gcactgtggc tatcanaaat tgcatacgtg cacatcntg gatcttaat acctatacat 240
acacaaacctt cataatgaat cttgactatc tacacaataa ggcgctacat ttcatgctgt 300
ctttcaagtg ttgtgactac ct 322

<210> 602
<211> 279
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 602

gatgatgacc gataacccaa agaatgattc gaagaatgag tcaacaagtt caagttcat 60
gagaaganat caagaagatt caagaatcaa gagaagttt atttcaagat tcatgagaag 120
atgaattcaa gttcaagag aagaaatcaa gaagacttca caagggagt attgaaaaga 180
tttcancaa acaaacatag cacagttctg tttcanaag agtttcttc acaatttcta 240
gtaccagagt tttactctct ggaatcgata ccagttcct 279

<210> 603
<211> 272
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 603

gtctcaagac tggactaata catntgctgt ccaagttta tggtcttgca ggtgaagatc 60
ctcataagca tcttatgaag ttccatattt tctgttccac catgaaggct cctgatgtcc 120
acgaagatca tatctttcta aaggctttc ctcattctct ggagggagtg gcanaagaat 180
ggctgtacta ctttgcttcc agttcatta ccagctgnga tgaccttaag agggtgttct 240
tggggaaatt ctccctaca tctatgacca ct 272

<210> 604
<211> 512
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 604

tgggnatga gactcctgat nacgtcttgg antttacgga cnctattcaa tactcaagct 60
tgtgaagtat gtatgttaat gaggaggagt ttaatgtggc attattggac ggatanacgg 120
ggtgtcttag taggatggtt gtgcggcct acacactcat taatattta taaaagtgtaa 180
aatgtatgc ttcaacattt aattctgaga tagagcacga agataatatc taactgatgg 240

cttaaattta ttgataagag ggagtagtct atagaaaagg tgtactttat acgatgaagt 300
gattttata aatgagataa tggcttagtg cggtctaattg tgataataaa gaatttgttg 360
tttgtgaaca ctattgttag gacattgggg atattttgg ttcttaact gtaaacgatt 420
acatgtaaat tttctattgc ttatatgata aggctgaatg aataggtgta aattaatatt 480
aattataatc ggagtttat gtctgtatct ag 512

<210> 605
<211> 394
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 605

ataatgatgg tccgagntat gttgtggagc ggntacgaac ccggaatggg tttaggcaaa 60
gacaacggcg gcataactag cctgataaat gccaaaggaa accgtggaa gtatgggtta 120
tgctataagc ccactcaggc aaatataaaag aggagcgttg ccggaaggaa gagcggaaagt 180
caaggctcgc ggatgagaca agaaggtgaa ggaaacccac cctgccaaat aagtaagagc 240
tttataagcg cgggtctgga ggacgaaggt caagtcgtca ctatatatga agatgatgct 300
ccgagtagcgc tgtatTTGgt acgaccatgc cttctgatt tacaactggg aaatcggcga 360
gaggaggaac gccccgactt ttacgcgaag agca 394

<210> 606
<211> 369
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 606

tttctctaca atngcatcac ctctcaatga gctggtaag aagaaatgtg gcattacctg 60
nggtgaagaa caagagcaag ctttgcttt gctcatagaa aagcttacta aggcacctgt 120
tctagctctt cctgactgtt ctaanacttt ttagctagaa tgtgatgcct ctggagtggg 180
agttggagct gtattgttac aaggtggca ccctattgtc ttatTTAGTG aanaacttca 240
tagtgccacc ctcaactacc ccacctatga taaagagctt tatgccttaa taagagccct 300
ccaaactcgt gaacattacc gttgttcaag gaatatgcat tcatagtgat catcaatcac 360

ttaagtaca

369

<210> 607
<211> 367
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 607

agcttgcc tc anagatgtcc aggaaggaca atgtctcaga aggaactagt tccgctccgg 60
agtatgatag tc acccgctt aggagcgcgg tacaccagca gcgcttcgaa gccatcaagg 120
ggtgtcg tt tctccggag cgacgcgtcc agtcaggga cgacgagtat actgatttcc 180
aggagggaaat agggcgccgg cgg tggcac cactggttac tcccattggcc aagtttgc 240
cagaaatagt ccttgagttt tatgccaatg ct tggccaa ac agaggaggc gtgcgtgaca 300
tgagatcctg gtttaggggt cagtggatcc cg ttcgatgc cgacgctatc ggccagctcc 360
tggata 367

<210> 608
<211> 280
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 608

cttatcaagt aaatggatca ttcttaacgt ccaacgcctt ataatgatca ccttcangt 60
aaaaagaatc gctt gattca cgcttaagaa agaactacat aggttcgatt tcctcatcga 120
tggagggtac gtatgagcaa aagccccgt attgtcgacc tcataatata aaaagacata 180
atagttaagg taatacatat tccacaattc taaaaaatat gttgttgtcc tttgagacaa 240
acgtgagagg tgctaatacc attctcaaac gtatatacaa 280

<210> 609
<211> 394
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 609

agcttgatt tcaaata ttttggtgcgc ttgtttaac atgttatgtt tgctactgat 60

tttaattct ttgaccctt gaatgaccaa attggcttc gatgtttca tgagacttg 120
agagaatttt atccttaca ttcaagcact ggtatcatgt tatttgacc attacaacat 180
aatcaatcct tanagcattg cagttntggtt atattgttag gacaaactga catctctatc 240
ttcatggtca gtttcttcca agatccaagc cttatggcc catgacttct ccataaaaga 300
tatatatatc tttctcttag ctntctacaa ccactgagat catccaaat tcactttgt 360
agctcaagta gtntcaaat tattgcacac atat 394

<210> 610
<211> 525
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 610

agtgatgnnn ttgaaaccc cagtaagtn ccnttgcttg tacgcatac tatacactac 60
tcaagtgcgc catgccnctt gatatacttg tggactcat ggtcaactatg aatgacaaat 120
tccttggat aaaggtagtg ttgtcatgtt ttcaaagccc gtactaaggc atacaaatcc 180
ttatcataag ttgaatagtt aaggtagga ccacttaact tttcaactaaa ataagcaatt 240
ggatggcctt cttgcaacaa cacaacccca atcccaacat ttgaagcatc acactcaatt 300
tcaaaagatt ttgaaagtt tggcaacgca agtatggggg cattagttt ctttgctta 360
agaacattga aagttcttc ttgtttctct ccccatttga aaccaacatt ttcttgagc 420
acttcattga gaggtgctgc caatgtgcta aaatccttca caaattgtcc ataanaattt 480
actaagccat gaacacttct caccttgtc acggacttan gtgag 525

<210> 611
<211> 386
<212> DNA
<213> Glycine max
<400> 611

atgcaagctt gcagatagat caatgtgagt caactttat ctttgatcaa attataaatg 60
tttgaattgt tctaaaatt ataataaaat caaatatgtat aaaataaaaa taaattctat 120
ttctgaaaaaa aaaagtcaat tctactttaa cctattgaat aaaattttaa taattcagaa 180

ttaattttt cactactgct aattcaaaca cacacttacc ataaacacgc gcgttgact 240
cgaaaatcaa ttgtctcccg ctccagcaaa atcaaattag taaagcgatt gccacataaa 300
tttatagtaa caaataacaa tcataatgc ctcaaagcta aaccccaaac catcttcttc 360
acctttctc tctgcacacg aatatg 386

<210> 612
<211> 273
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 612

ctatatgata tagtgttattc tctatattat gttcgatgt ggaggaggct aaattatcat 60
. tctaattctc ttatagtnt aattgtgtaa tcttgattgt ataaattatt aaatatataa 120
acatttgttc ttattttatt atactatata gattgtctt acattattgt atatcattta 180
aatattatga ggatatgaaa ttataattta accttataa aaatagatgt aacgcaacat 240
agagactgat gctactttga tattccaatt gat 273

<210> 613
<211> 273
<212> DNA
<213> Glycine max

<400> 613

tatcttgtc attcaatatc ctgatgaggg tgttcatat gttctcaaga ctggacaaat 60
acattggctg ccccagttc atggacttgc aagcgaagat cctcatacac atcttagcga 120
ggttaatagt gattgatcca ccatgaagcc ccctaattttt caggaagatc atatcttact 180
aaagtattta ctcattctct ggaaggagag agaaaagaat gcgtgttcta cattgctgat 240
agatccatct ccaactggga tgaccataag aga 273

<210> 614
<211> 353
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 614

gagatcatcc nctcaacaac attatgggtg atatctcana tgggttaaca tctagacact 60
ctcttaaaga tntatgcaat aatatggctt ttgttatccat gattgaacac taaaatataa 120
aagaaggccat aatagatgt aattggatca ttgccatgca agaagaatta aaccaatttg 180
aaagaataa tgtgtggaaa ttagtagaac aacctggaaa ttatcctatc atatgaacaa 240
aatgtgttt tagaaataaa ttatatgaac atggtataat tattagaaat aaagccaggt 300
tagtagcaca agggataat caagaagaac gaatagacta tgaagaaaca tat 353

<210> 615
<211> 122
<212> DNA
<213> Glycine max

<400> 615

tgctattgta acatccactt tattagcaat tgtggatag attagaaggc cattgatgg 60
gacgctaata ttgaagaagg aacaattcta ttatagaaac acaaggggaa ccattttac 120
ac 122

<210> 616
<211> 345
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 616

cagtgtttat attgcggttc caatgtggcc agagggttac ccagaacatg gaacggttca 60
agcaatgatg gattggcaga ggagaacaat ggatatgatg tacaaggatg atgctggagc 120
actaaaaaggc aagggtaatg aggaagatcc tctcaactat tcgacattct tctgcctagn 180
taatagggag ctgaacaaag aatgagagta tgtccccca gaaagaccag atcctcatac 240
agattatatg agagcacaag tgtcccgacg ctttatgatt tatgttcatg ccaagatgt 300
gatacgcatt tctgttaatt aagttacca catcttccaa gatac 345

<210> 617
<211> 288
<212> DNA
<213> Glycine max

<223> unsure at all n locations

<400> 617

ttctaacaca ggtattttca cggatttgta ctggatgct cttatactan gaagtctctg 60
cttataatant ttactttaga ttaaggagga cagatactaa tgctcccttt aacttgaat 120
atccttcctt cttcatcg agagcaagaa aaacttacta atggatggaa gcatctatac 180
catgttgcat ntacaatana agcaaaaacac acaatctcaa tataacttct attcaatagt 240
tggccaaaga cttcacaaat gagaagagat tagatattaa tataaaga 288

<210> 618

<211> 459

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 618

ngcaagctnt ctaaatgttt cttagccag tctcaggtag agtacttagg acatttggtt 60
tcgcacatcggn gagtggagcc cttagcatca aaggtcactg caattcagca atggccaatt 120
cctcgtacaa cgaaggccgt aaggagctc ctgggccttg ctggctttt tcgcaggttc 180
attcagaatt atgccattgg tgtggccccca ttagtcaaag ccacgaccaa agaacccttg 240
cattggacat ctgagacaca tgaagcctt gacacttga aacatgcctt gtcaatagct 300
ccgggtgttag cttagccaga cttcaacctt cccttcacag tcgagacaga tgcgtcagga 360
gttggtatgg gtgccattct ttcacagcga ggccacccca tagcatttt cagcaaacct 420
tttagtgcca agtactctga tcataacata catgcgaga 459

<210> 619

<211> 492

<212> DNA

<213> Glycine max

<400> 619

cggaaatgata cactttacta tctataatct cagctctcag gagctgagct agttattaaa 60
gggggtgttg tagcttatct cttagattctc aagaaagttc tctcacatat tttctcaat 120
ataacttctc aatgaaccta cctagtcata gaatagaagc atgtgtcaca cttgttgg 180
ctatgatgaa tgatactttt atgagacaca ctacacagtt ccacttgtct tcctctacta 240
taacttaaac tgaatctacc cctgctcttt ttctttccct acatctaagc atactctata 300

tgcttcttat ccaagacatc actcttcgt gtattccctct tctttcatga gctacataacc 360
tatcggtctg tccaggccat atctatTTT ctaaacactt tgtgcttcta tagttccaac 420
atTTCCCTTC tggctgctct tactcttTC atactgtctg ctgtggcTT taatgttcct 480
tttccttac cg 492

<210> 620
<211> 416
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 620

ctaagaagaa agtagaaaaac atatcacctt tctatTTAac tatctttgt attgtatgct 60
tggcaaatct gaatggtaact ttccTCTGTT tggggTggaa cttatggatt tccatttccc 120
aaaagggtga ctcttctgtc acttccatct tcataatgatt ttggatgtg gcttcagttt 180
tagccattat gtggntgctg tggcgccca attgttgac ttcaatgata aattgtgttt 240
cttactcaac atcaattcat tggaaattaa tgcttcTTT aactcagtTC tatttgcag 300
aactagagaa tactatacac gtgtttctgc atgaacacca gctcatgcag ctacttcatt 360
aacgcataata taattaaagt tggctctcag aatntctcca tatcaccaac tatgat 416

<210> 621
<211> 328
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 621

atagccccag gactagagta cttgcataag ggatgcaaca ctcgaanttt acattnTGAC 60
atanagccac aacattcttt nggatgagaa gttctgcctc aagatatctg attntgggct 120
agcaaaggct tgccttagaa atgaaagtat tatttccagg tctgatGCCA gaggaacatt 180
atggatgtt gctccagaaa atttggcaga atttcacaca natctgatgt aatcttccac 240
tttanattta aaccacctaa accttaatgg gtAAAATTAA ttcttattatg cattanatgc 300
atcttatctt tgacttgaac tctacaat 328

<210> 622
<211> 439
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 622

ttatagacga atgtttcaact tttcttccac ttctaccgag agacataaaa ttttcaacgt 60
ctttcatagc atcaaactga aaatctctaa ctgcccttcc aatactaattc gctgcctcaa 120
caattccctt ttctcccatt agcttgctcc tgtttagtga cacaatgcga atgacaatgc 180
agtttccaaa gtacgttgaa ggaatcgaaa attcgggacg gttacggcaa tctcccgaaa 240
acttcaagct gtaaatttca tcattattat tattatgc aatagtgata gcattgaccc 300
cttcttctga ttgaaccttg caaaccacaca tcanagcact tgtcataaca aagggttgcata 360
tgtgcaatga ctctaattct aaccctataag tacttctaca ttcaatcgac acccatttct 420
tttagcttcgc aacatggcc 439

<210> 623
<211> 385
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 623

taaacctctt aatttagtca atgcaatttc ttttttatcc cttgttctaa ttacatgcat 60
aatgcacatctg gttntccctag ttttcaatt gttaaaatta ggtccatgtat ctaatgtatt 120
ttttgacatg atagttggctt gtctgacttg ccacatgata canagcaatg gcatgagttc 180
actggttata gccactgttt tgtctgtcat tgtaagaatg nngaaattgt gaatatgaat 240
aggatttgaa caagaatctg ttcttgact annaaatatt attttatga cacatttan 300
aacggctctaa tcgcttanaa tgtagcttg tcaatttata ataatanaag tatataatga 360
caatttctaa atgtcttaca tggta 385

<210> 624
<211> 335
<212> DNA
<213> Glycine max

<223> unsure at all n locations

<400> 624

gtgtgcgtt agttggtaa gggtggtca cgttgtgatt ggcggaagtg gttggttcaa 60
ctccttctac cattggtgag ggaagttgaa tattacattt taaaaaaatc ttacatagtg 120
tcgccttcc ctgtgaaata ttnttcgcaa tagaaaactaa tcttctgcta gaattatgag 180
tggaaaactaa ccaattaaat atttatacaa ttaatactt cagtatttct cataattaaa 240
taaatcatga tgatcaaata tatattctct tctaaaagag aatcaattca acatacacaa 300
gtctacagga aaactatatt atgagtaaac tatca 335

<210> 625

<211> 514

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 625

ntganncccc tggtaaaccc tatagttatnt accngtcaact atttcgtaact caagcttatg 60
cttctaccgg ttntggacta ccaaattgtc aagttgttt tgcataaca aaacaatcat 120
attcatctca caaaaaaaaaa tcgattccaa atcgatgtt ttggctaaag ataatgacat 180
ctcttaactc cttggcaatt gtatcaattc gttcaagtaa tagtttaaac gataaaaatcg 240
agtgtgagtc cacaagaact ttgactgtac tcagagttt tatatatcca attttaagta 300
ataaaattaat tgaattgaat atttgttgag tgcgtacgca taaatataat tttgatctaa 360
attaaactac aaaacanagc atgtcaagg gtgagaaaac aaacactcaa aacagtgaag 420
taacgattga tgcagaatta tgaacatgtt gggctcagt cagcctacca gaactactct 480
ngatgcaaca ttaaggatnt ntctctattt aacg 514

<210> 626

<211> 314

<212> DNA

<213> Glycine max

<400> 626

gacgaagcgt gttgtgaatg gcgtctcgcg tggctagcat ggtatcgaca gcatcattgc 60
gtgatgacca cttgtatgtaa ttggagatgg ccggaggagg cttaccatat gtgacacctg 120
atggagtgaa gtcgggtccg gagtgctgag acgtgttgta agaccattca gctagggcta 180

agaattaaac cagtatgccg gggtgtggtg aacgaaggaa cgaagatatt gctcaatcg 240
accgctcatac acttcggttt gccccatcgga ctggtgatga tacgcccgtac tcatgcgtat 300
cgtcgtccca ctga 314

<210> 627
<211> 264
<212> DNA
<213> Glycine max

<400> 627.

agctgttagga cgtgaaatca tgtgcagtca tgtatcttat agtcctctca cggggggggag 60
gttgtgccccat gctctcagaa tgtgcaaaat cagaatgctc agaatcagaa tcctcagaat 120
cagaacgctt aagattatta cgcttccaat cgagatggtc atgagcacca ataacatact 180
gcacagattc atcatgagcg gcctgctccg gatgaccaaa aggaataaca tgatgcctaa 240
ctcatctatg aaatgtccta tcta 264

<210> 628
<211> 522
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 628

tggggaatga tacctnagtg agtangcgtg ctattacctg tcactatcga atactcaagc 60
tgggagagga tgcttcaatg gaggaagaga ttatggaca gatttataga gggggtagca 120
cggaaattgaa ggaataatat acggagagaa gtggAACCTT gaagtatgtc ttacatgact 180
ctcattccatc aaagttacaa catgtgttgc acatgcttct cttagatagac taggttagctg 240
gcttgagaag ctttgggttgaaaactgtct tgataagctc atttgtgaag cattacgtgc 300
gatgctagag cttagctcca cacacccggg tgataactaa actcacctcc tggagaagct 360
cccttaacat gactcctaac gaagctatag cttaaactaca catacctccc taataactaa 420
gctcaccttc ttgtaatgag aagactttaa ctgagctacn tggccctgt ttagctata 480
ctcacacactg tgacaaataa gcaatggtat gcaatacact cg 522

<210> 629

<211> 396
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 629

gactcattat ccntracagn tatttaaca gaatgtcttc ctggcttact ctgatttctt 60
gaaggagaat ccccaataa atcactcacc ttttcttcta cgagagaga acttctcggt 120
ggagcaatct tcttgatgc tttaggtta gaatccttt tcctccatcc accaaaccat 180
cctttctttt cttgcttatt ttctccactt ttgttacaat ttccatcttc aataggaatt 240
tctctttctt cataacagct gtgacgatgc ccaatgatca cctcatcact ctcattgttc 300
aagtctggcg actccaacct tagtgcacatc tccaattgcc ttcttcatc ctcggncaaa 360
atatcattga gctttcact ctcttggta ttttca 396

<210> 630
<211> 294
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 630

tcttacacgt gtaacacaca acttcagctt agtggttaa gactctaagt gtagcgtcca 60
tggaaaggcga agcatccata anagttacta tccatagtgc tctatggcct tcaataccat 120
caccatcaag caagggagat gcttcaacct ccaacccagc tgctgcanaa tggccactca 180
atatacaaga atgtattcct catacaatag caccaccaca tttagaatgct cgagttgatc 240
caaaaatgca caacatataa naatgctaac aataaaagtc ctaaccctta tcat 294

<210> 631
<211> 365
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 631

tctcttatgt tgcaagagtct atgaattcta gaataatata acttcaacca ctanaaatga 60
cttcaaattt tatttaacat ctaagccttg tcagctcana tggatcaaattt cctcaatgtg 120
aaagggngaa cagactcttc tatttataag agtcttanag gtaatttagtg atttacaaat 180

atcatagcan ataattaacc ttcttaagagc tntccgcctt ttagcanagg ctgctttaga 240
cctgatcgat gaggtacaac tcccttaaca aggttcgtc aatgtggtcg attgattatg 300
gagttaattt tcaacaactt tgtaggata tgcatacatac atggatgatc atcatacaca 360
tacta 365

<210> 632
<211> 396
<212> DNA
<213> Glycine max

<400> 632

tagttttgg ttttggtgaa cacaattaat ctattaccat gtaactgtaa tcgattactt 60
gggtgttgt gcgtgatgta atatgttaca tctctatgtc ttcacccgt caaccactat 120
tattactaca ttacttccg ccacccaccc cagccacta aaatgcctca gcctctctt 180
ataagccaac cttaccctat gcacaaatca caccaacacc tgttctaaac ctcctgttac 240
caccccttc tgaatacataaa tatcttatgc tctcaaacc tctatctcct cttcaagatg 300
acaaacaatc cttaagagc aaataatatg gaaaaatatt ccaagaagaa gcaaggagaa 360
tgttcatagg atgttatcaa cccgttggac ttggaa 396

<210> 633
<211> 517
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 633

gcgggaatga acccatgtga naccntgttg atacgatggc atctacgtga cactgtgtag 60
tgctccagcc cgtgcgatta tgaggtaccc gtctcctgtg gtacttaggtg gtattcgggc 120
gagggtgcat aacatttctg acgatccacg gatgacctat aagcacccca tacgcttagtg 180
gcccctatg gcagagctca ctatctcaca catgacacac atacgtggtg gtgtaagcac 240
cgggtacacc tcaaggccttc tgagcttca catcatacaa gtaattcaac aattattgtt 300
ccgaactcac acatgatgag ctgaaggcgt aggcgcagaa ctctgctgga acacatacca 360
gtctcccgac tttcacatg caaatatccc atatgcattt gctatggtct acgacgttga 420

ccgtgtggat cgactagcat agactactgt gcgtatctacatattct acatcttgc 480

cgtatgtata tgctatagaa tggctagact cacgtag 517

<210> 634

<211> 390

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 634

ctttagctat aattcattgt ttgtgttatg agtctacatc aaacaattgt attattgt 60

tttctgtcac aatcaagtga tagttatgt ctccatatgt gcgtacactg tgattatgtt 120

ttcgtttcta aaattcattn ggagtatcta ctgttgattc tagatgagtg atccttctt 180

atttacaatt attgtctcct aatcaatcga gtgttcatct tattattggc tgctcatatt 240

ccaatcatgt ctatgtaaaa tgcttgataa tctttcttgg tgttacttct aatacgaata 300

aagagagact tcgccttga caatcacgtt aagatgtttn gagaggaaat acttgagtag 360

acatgatcat tgtacctata gacgaaagtc 390

<210> 635

<211> 252

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 635

gattgatagt tcactaagaa taatcttata aagaataacct ttccttttag caatgaaaaa 60

gtgagtcaccatccccc attttgtnt ggacaacaca cccttcttgc tttaaggaaa catcaagtcc 120

attgtcacat aattgactta tgctaaagcan nnatatgtta agctctttaa caaaaagtac 180

attatcaatg ggaggatagg gatcaatact tantcttca tactccaact atttcctt 240

ctatccctt cg 252

<210> 636

<211> 393

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 636

cagctccatt gatatacattt atttaatgca atgcacatgt ttcgaataag atttcttgta 60
tcacaggaaa agatctaattc caagttgaat tacggatcat actgattgtg gtgcgaaaac 120
tggcttgaag tagccatctg ggtaatatcc aaaccacgaa gtttcctctg gtattaaaac 180
agtgtcgccc tcacactgtt aatgtataaa ataagactac aaatatataa gcttcctca 240
tgaaatcaat tagcaatatt ctattcataa tatcacaata atacatttag aggacttacc 300
atgataagta ccanattctg caagctactc aatctttct tgtaagaagc atttctctta 360
tctggtattt cattgccatg cactggaaga aat 393

<210> 637
<211> 391
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 637

gctgtctcta gatagacaat cttgccctcc tgaccttacg gtgacttacc actttgatnt 60
cttttttgt tggtagttaa ctgtaccatt aaatataagg atcatataag aaactagctc 120
acaagaaaaaa tctggtaaag actacatatg annaaatagc gtggcttaa acaacaattg 180
taacgtaata nactgaaccg ctggtaatt tcttgaataa ataaattcag atcttcacaa 240
tgataatttc ataaactctt actgtacaac acattgactg aagacaaaca ttatgcataa 300
aactatagaa gcatggcata tggcatctg caaccaacag gaatgtatgt cacacattga 360
aacagatatg tnntatctca tactcattct t 391

<210> 638
<211> 416
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 638

ctgcacaagg ctcttaatat ttgaagagta tccttgagga acttcaccc gacgaagaca 60
ctaacaaaaaa cttatcttct cttcttgga caaagtatgg caggctgggg gcaagtaaat 120
tttcttccca tcagaccttg tatgcaactg tgatcgtata cccatatcag cttagatctg 180
acgggtattc aagccatcct tcgtcttgcc ttgaatgtta aggagcgtcc caatgactct 240

atcacagaca ttttctcca catgcataac atcaatacaa tgtctaacct caagatcaca 300
ccaatacgga agatcaaaga atatggacct cttcttccat atgcaactnt gactattatc 360
cttctttga gtcttcccag atacagtatt cacgtgttca acccgataat atacct 416

<210> 639
<211> 298
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 639

gttcttgacc atattgaacc ccgggtgacaa caattgcttc aaggacagag acaatgaaga 60
ggtcgctgga caacaggatc aacaaatgac ctatagtatg gttcttgact tagatcgaac 120
ctgaggacaa cgatattaac caacagacca cacaacaaat gaataacagg acaagagaca 180
atggagctgc acaacatagg gttaagagag acaaatgatg agggataat ctcanaatacc 240
ttgcaagcat ggttagaaact ccaaacccaa actgagcata ctcaaattat ccaatctt 298

<210> 640
<211> 405
<212> DNA
<213> Glycine max

<400> 640

atcactatac acatttagatc tgagtaagat ctactattga agtactaaaa tgtacacaca 60
catacctcgc cacaaagagc aatatacttg gaaaggagct tggattct atcctgctca 120
tcttcctga gaattcgtag ttctcctca ctcacagatc gtgcaagaag attcccaact 180
gcaacaaaat cacaaaacta agctaaaacc aataaaattt caatgcacac acacaatcga 240
aatttagagct gaaacataac tccaaacttac atggtgtata gacataatcc ttggtaat 300
tgtgtgtgac atggagaaca gcaatggaaa ttggatgaga ctccacttct tgagagccca 360
attcatagtc ttaagtccat cttgcacatc attaccaaca gcaac 405

<210> 641
<211> 312
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 641

taattaagca atatggatga gatatctctc taattcagga tctaattata acatgatatg 60
atcaaatgt aagaactaaca tcactcaatt aagatatggg aattaaaatt acacagtatg 120
tgaatataaa gaaactcana ttgtatctaa actttctttt ctactaatat tactattaaa 180
taatatgttt taactatgcc tttaatgaaa gtgtgtttt actatatatt accaaagaaa 240
catagtcaa cctatataaa acgtgtatnc gaacaaacct actatcagta agtataacttt 300
atacgtaat ct 312

<210> 642
<211> 482
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 642

tatgaccct ganncccttga tccctgatta ctgcggntta agataaccgc angccgaagg 60
tgtatatgtat ttatgaactg gccatcatttgc tccttatgca agtctngcac ctatgttgc 120
tggtcgacac aaccatcata ttccgactga gagtgtgtgc ttaaagaaca tcacgtcgaa 180
tcggccgtat cgctgtcacc gctcatatag agtaacatta tgtggactac ttatagcggc 240
gattacaggt ggggtgtttt ttcatatatac acatgctaca tcccttctc ctaaatgcaa 300
aagacttcgc tcgcttcata gcttatactt acttcctttt gaangacaga cacatcacat 360
tgaacatcac caaaaatatac tttggcagg ctgccaagac aaagacttat gctgtatgg 420
atgaccataa cttaataggt cactccccg agacctagag aaaccgatct tcgacttaaa 480
cc 482

<210> 643
<211> 514
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 643

tggggatgac nccattagan ancncgttat ttgcngacac tatacagttac tcaagctntg 60
agaaaattatg gctgaggcta agacatgttgc tggcccatt gttcaaattgg aggcatttat 120

ttatTTATTT ttaatttatac atttgggtaa atgtcaaatt aatatggat aagtcattat 180
agatgttgta acttttgaat cgaaagtatt tttttttaa tttatgactt taagatcgta 240
ggagttttt tttagtactc taaagtcgta aagtctattc gttaaatttt ttatTTAATT 300
cataaatgaa tgtacaactt caatTTTTT attcTTTTC ataaaattgt aaatcatttt 360
ttaattcaat tatttaataa ataaatgtat ttttactttt cttagTTTA tataatattt 420
tctacatatt ttttatatag acttatctaa tatatttcct atactagttt attcaatatt 480
gtcttaatcg ctggtatatt atcttatttt atat 514

<210> 644
<211> 309
<212> DNA
<213> Glycine max

<400> 644

agcttgataa caatgtctct tctgtttagc attattaacg gcatgctcct cttccatttt 60
gcattcctgct acaggaaggt gccattcggt agagctaaga attcttaca gggctgatgg 120
ggTTTatcaa caaggttagtc ccaacttagga tcggacactt ccaccgttgg atgataggca 180
tcttctacaa tattcTTTat ttccatgggta tcttcattct cttatatttgcgcattcctct 240
ttggcgctct gcttaaagag ccagaaccat tacccattgt ctctaaaat gcgtgaatct 300
ggggcattt 309

<210> 645
<211> 310
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 645

agagcactcg atctatataa agaactttt cttcaacaac aacacccat canaactcat 60
gagaataat aatgcanaac taacactaat aagatgtgc acaaataccc attatgggtaa 120
aactcaaaga aagaagaana aggcttacca tccaatagtgggtcacaag ctcaaggaaa 180
tagatgagtg caaattgtcc taaaggaaa taagccat tcttgagagt gaatgaaaaaa 240
ccttcttatac gttggaggag aaaatggaa agctctgaga aatgagtaaa ggtgcatagt 300

tncaaagtat

310

<210> 646
<211> 297
<212> DNA
<213> Glycine max

<400> 646

agcttatcga gatccgtat ggataggcaa tggttcgca taatgtggaa cctgacgttg 60
ttgatgtaca catgcaaatac ggagaagatg gagagttatt tgatggcaaa cacaatcatg 120
ttgttcagct tcacgacttc cgccgtgcatac accactaagg tgaggatgtt gaagccattg 180
ttgcggataa gaagcatcac gtcaatcaac atcaagaata ctttcgcgaa ggtgttgttc 240
ttgttccaat gaagcgtatc agaggtaaa gtgatgcaat ggcgaagtgg gttcgag 297

<210> 647
<211> 372
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 647

ctaactgngc agtantaacaa aaatgtgagt gacatcacct tgagggtggc aacaaattaa 60
gtgccttcac cataatagca cctgcacactg aacctatgaa gtttcattct acacactgaga 120
aagatgatga gatatttctt cataagagaa agttccaac gcacatatta cgcaactgac 180
cctttttcc tctcatgact taaccacccc tcctacactt attattccca ataccatatc 240
tcactcaccc gtacatcacc attctgcacc taccttacct ctccctgcag acaaactctt 300
tatcgcccttc acatgtaacc ccgtaaacac tacaatctct aatatatgat tgctcaactc 360
tcttcaacct ct 372

<210> 648
<211> 375
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 648

ttttagccgt cgacgacccng ngatccttag agtcacactga ggctgctgct tacatcaaca 60

cttcagggc tgtactactt acatggattt atgggcstat gcagttgaaa gcctggagaa 120
agaggatgcc tatngttgtt gggatgaatt ctccagattt acctggtaa actctatcg 180
agagaatcaa aaccttgagt attcaagagc tgagtctaag acttcaaaga gagaaagact 240
gtgtcatcaa gagaatcagg agtgaccatg gcagagaatt tgaaacagca ggttcactga 300
atttgcacat ctgaggcata ctcatgagtt tttgcagcat tacacaaaca gaatggata 360
gtgagaggaa aacag 375

<210> 649
<211> 234
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 649

tttgagcgtc tcgatatatg acgagactac atcatacatc ttagtaanaa gttatagtcg 60
tttgaatatg ctcagagctt caacattcaa ttacgagcat ctcgctatat tacggactc 120
aatcagacat ccgagtaaac agtttgggt ttgaattgtc tgagaggact cacattcaca 180
ttctagcgcc tcgatatata tatggactct atcacacatc cgagtataaa gtta 234

<210> 650
<211> 407
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 650

tgcaagcttc tttgagaaaaa cttccttgag aagctagtgc ttagctacac acacccctct 60
cataactaag ctcacctcct tgagaagctt ctctaaag aagctagagc 120
ttagctacac atacctctct aatagctaag ctcacctcct tgagatgaga agctagagct 180
tagctacaca cccnctataa tagctaagct cacccccatg aaaaaaaaaa catgacaata 240
aaaaaaaaaa agtccttact acaaagacta ctcaaaatgc cccgaaatac aaggctaaaa 300
ccctatacta ttagaatggc caaaatacaa ggcccaaacg aagaaaaaac ctattctaat 360
atttacaaag ataagcgggt catgcttagc ccatgggctc gaaatct 407

<210> 651

<211> 337
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 651

cgccttaacta aatatgatat tgagtgttt ttacatgcat ntacttaaca agaaaacttan 60
acattagnta agtttaaact atatggttgg aaaaagaaca agtggaagag tgaaggtttc 120
caaacaacaa tccttgacaa aaaataaaaa aaagctggtt ctagcaaata aatcatatca 180
cactaagaat gagaataat accagcatcc ctatccaaaa aatattcaag gatgagacgt 240
gtaaaaggtc acgaatttca tgctgcta at gatatactaa acantaataa tctngtcctt 300
ttataccctt tanggttata tggagatgga cggag 337

<210> 652
<211> 178
<212> DNA
<213> Glycine max

<400> 652

cccatcatga tgctctttct gaacagaaaag tacctgtgaa cgtgcactaa attgcctatc 60
aactcaactac gagagaaaatt gaaccttatct taaactctgt gatatgagta tcattacaat 120
tatgaggcat ctcagtgcataaaatggatc aggttggtcc ttatgacctt gctgccta 178

<210> 653
<211> 354
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 653

tagtacaca cacccatcta anaactaagc tcacccctt gagaaaacttc ctgagaagc 60
tagaagcttag ctacacancc ctataatagc taagctcacc cncatgacaa anaaacatga 120
naatacanaa aanatcctac tacaaagact actcanaatg ccctgaaata caaggctaan 180
accctataact actagaatgg cccaaataca aggcccagac gaagganata cctattctaa 240
tatntacaaa gataagcggtt ctcatactta gcccatgggc tcgaaatcta ccctaaggct 300
catgagaacn ctanggtctt cctttggatc tctagccaa tctacttgga gtct 354

<210> 654
<211> 445
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 654

taataggta agcactacaa catttataat ttcaaagcaa agcacataat aatagctatt 60
gtacataggt ttgatcacac ataacacact atagaatatt attcgtcata attaacaac 120
aattctaaaa gctatagcgt catggtaactt taagatgaca tatagagttt taggtttatc 180
aaagaatttc ccataacaaa tccaggacta tatcccaaca ttgcaaatacg tacaagagca 240
agcaatcaat atactttaga atctcacacc catattccat ttcaagcatt atatttttc 300
atattnagac ataacttggc ttatgttcta ggccaatatt ggactttatt ttacaccagc 360
ttattggact ntaagaatac atcccgaac atcaaatacg tcaaataccat tactgcaaaa 420
tacaaatacc cagaggaact acaac 445

<210> 655
<211> 520
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 655

atgttaattga aacctttgtta agnccgtca ctatagaaca ctcaagcttc taagattatt 60
aataatatat ccattattca atatttcaa ttaattataa aaacaaaatt atttatctaa 120
aaaagaanaa ttaaaaaaaaat attttgaaaa caaaataatt taaaattact aagagaaaga 180
gcaactaaga ttttgacaga aaaaatgaat gcaaaaataa cacaattaaa attaaaaaaat 240
aataaccatt aatgtcttac attttatgc ataaacatat atattacttt taatttaaaa 300
ataaaaaatat tttagtcatt tgtgtgaaat taaattactt acaacaaata aatttaattc 360
aattctttaa tagaaaaact ctttatatat atatatatat atatatatat atatatatat 420
atatatatat atatatatat atatatatat atatatacac acgtatcagg gacatatgtg 480
ggataatatg actacgttat atatgcgagt ggagatgctg 520

<210> 656

<211> 420
<212> DNA
<213> Glycine max

<223> unsure at all n locations

<400> 656

agtacaataa gctntatgtt ctaaatctac tggaagatga catcccttgc catgcaccat 60
ctagaatggg gaaagactgg taggggtctt ataggctatt ctatacaccc acaaagcatc 120
atccaattta gcaaaccaat cttaactaga attctctta tctttgtaaa tgagacttca 180
acttgaccat tttttgagg gtggtaaggc gatgcaacct tctgtctgac attatagtgc 240
tccaaatgcct tctgttagttg cacattgcaa aaatgggaac ccccattact gatgagaact 300
ctaggagttt ccaacccacg aaaaatattc ctttcacaa accgaatgac tattttgca 360
tcaccccttct gtgtggcaat ggcttacacc cattttgata gaaactcttt aaatgacaca 420

<210> 657
<211> 265
<212> DNA
<213> Glycine max

<223> unsure at all n locations

<400> 657

gtatccattc agagaaatnt cctttggga ggtcatcatg aggccaacaa gattccttgt 60
gtgaagtng acacagtttgccttctaann aataaagggg gccttngat taaagatttg 120
tctaaattta atgaggctnt acttgacana atgggtggg agctggctaa taattagaac 180
caactntggg caagaatctt aatctccaaa tatggtggtt ggaaggagtt gatctctgg 240
ggaaagagca nattntcctc tcata 265

<210> 658
<211> 360
<212> DNA
<213> Glycine max

<223> unsure at all n locations

<400> 658

tgaccttggtt tacagcgacg ggcactctt agaactgaca gaggcccata atcagagctg 60
gaaacgccag gaccctatttgcacttctcg agtccactgg gtgtctttagt ggtgcgatcc 120
ctacaaacaa tagatgacat caganatcg ttgagcgatg tgcataactta cctatgtcac 180

gatggcatga ccttgctggg ggcacggca ccctgttaga ctgacagagg cccgttaacct 240
gagctggaaa cccttagggcc ttgttgggtt tctctgagac cacggcgtgt ctgtggcg 300
cgatccctag caatagtgg aatcaactga accatatgca tacttaccta 360

<210> 659
<211> 360
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 659

ttatctagta cattgtnttt gaaacaagca ccctacagta gaattgccaa caatagacat 60
tgtttcagtt agggtttca attatcatta ttgacgactt ttagtgagcc tcctaaagaa 120
tgtaaacgcg tggcttatgc caccctcttg ataaactaag aagaagaagg tgaataataa 180
ataatctttt ctattaaaat aatacgttag ttggtagaag gtatttataa cattaaatag 240
tactctcttc atttgtaaac gatattttn tatatacaca tattaagaac gtcaaaggat 300
atataattaca taataagggt tcattgagtg ctatagattt taatattact ttagtttat 360

<210> 660
<211> 390
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 660

agctttagg gttatagtct cacgattgtc acgtgctcat gcaacaattt ttagccgtgg 60
ctatacgaga catcttgcca aacaaagtca gggtcacgt aacttgcttg tgcttttct 120
tacatgtat gttagcaaa gtgattgtc cagtaatgtt ttagtgatgtt gaaaacgaga 180
ccgcaattat actatgccag ttggagatgt atttcccccc tgctttctt gacatcatga 240
ttcacttgat tgtgcacatgt gtcagagaaaa tcaaattgtcg cggccctgtt tatctaccc 300
ggatgtaccc ggctgagcga tacatgaaca tctttnaga gtatacaaag aatctatatc 360
atccgaaagc atctattgtt gagaggtaca 390

<210> 661
<211> 111

<212> DNA
 <213> Glycine max
 <400> 661
 agaatgaagt ccactcaaac ctgaaatctc caacttccac tcgttagacac gcacttcacg 60
 actaccgaaa tgccctcctt ttgcgatctg gagcgaaat gatggccaaa g 111

 <210> 662
 <211> 404
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 662
 agcttgtacc agccactana cttcaattt caatttgang ctntccata ttcttgacct 60
 tcctcctacg agagaccaac ttcaactcag tatgttcctg tanataatta taattaatta 120
 gaataaataa aattgtatat aaaatntaaa tatgttataa ttaaagaaat aactacctct 180
 cttgcccaca cttggctac cacatgatta acatatgatg tcaacactaa tgtatctgn 240
 ggccccctg gaaaacccta tgaatcaaca cctacatcct ttgtaattgg atcatgaggt 300
 tcctcatgag tctcatcagc agcatcatcg atatgccat tatttcgac aatagttgca 360
 agtggcatt atctacgtgt cgacacccat aatcttcgac gctg 404

 <210> 663
 <211> 319
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 663
 gcacgagaca tcagggtta gtattccaag atacaacata tatcgcatga acattgttag 60
 atagagaaca tccttaataa catcagtcat ccagtaggaa gaccaacacg ttctttatct 120
 gtcttcatac accactactc acgtgattta atttggata gtttagttgc atacttgtcc 180
 ataccacgca ccanactntc atccaaaggc acttatttac tgaaccacag ctntaccaag 240
 tacaacagaa tgctcggag ttggatactc agtattcact taccggttta tactactttc 300
 gtgatccagt gcacttgtc 319

<210> 664
<211> 349
<212> DNA
<213> Glycine max

<400> 664

aagcttctat agaatgttcg ttcctaattt ctctacaattt gcatcaccc tcataatgagct 60
ggtaagaac gatgtggcat ttacctgagg taaaaaaca gagcttagcct ttggtttgat 120
caaagaaaag cttactaagg cacatgttct aactcttctt gactttctaa aacttttga 180
gctacaatga gatgcctttg gagcgggagt tggagctgta ttgatacaag gcgggcacct 240
tatttcttat attagtgaaa gacttcatag tgccaccctc aactacccca cctatgataa 300
agagctatat gccttaataa gagccctcca aacttggaa cattacctt 349

<210> 665
<211> 295
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 665

gtcacattgt caaagctcca atcttgnnt taaaatagag agagcaacaa cagcaanac 60
tacaatcgaa gaaaataaat cataatcgaa gagcanaaaa aatatcagaa acaaggttct 120
aacgtttctc tcacacaang cttttatctt cctctnctaa tccatttttc ttttctttg 180
gtattccacc anagacaatt ttttctaat ggaaaaaaca ctcgactggc agagaaatag 240
tgaagtgaaa agagagactg agagaanaga tattattctg gtgacgctga tgtgt 295

<210> 666
<211> 429
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 666

agcttggcag tgtgacattn tcttctgagt attaatgca gcagattaat gagttttcat 60
tcttatact gaatttccca tcgaagcaca aacacaatca ggagcactag gtttaatata 120
atccttagct tgagatgccaa attcctctg ctccctgcgt agttatatgt caaaagtgg 180
aaagactctg catcatcata acaatcaaac caaataatgt gcataaggat ttagctttc 240

tagagtgagc aagtgtgtaa agtaagaaaag taaaagtcaa tgtactgtt atttgaattg 300
atgtacacctgg tggccctaac taactttaaa acaatgggaa aacaccctta gtttctca 360
ttatacacattt actaacttta gaggagtcaa atccctgcat cagaatatga acaagaaaac 420
atcacctta 429

<210> 667
<211> 459
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 667

ntatcaccag gatgatcaaa atgttttca tcattgcttg agttgaaagg ttggaaatgg 60
ggataaagtc agcttttggaa aagataagtg gtttagggaa ggtcctactc tacaacagaa 120
atacaatcag ctgtttctaa ttaatagaca gcaatctgac cttatttcaa tggatggctaa 180
tttctctcat gatacatgga gatggactt gaaatggaga aggaacctgt ttgaccatga 240
aagtgatcta gctgtcgatt ttatgaaaga aataagctct tttcatttc agagaaatgt 300
taaagacatc atgacctgga aagctgatcc tagtggtgta tattccacga ggtcagcata 360
caaattcatg ataaccccct cttcccaagc ctttgatctg agatcctcaa ctntattatg 420
gaaattgaaag attccccaga agctgcagtt ttcaacttgg 459

<210> 668
<211> 306
<212> DNA
<213> Glycine max

<400> 668

ttgctgtgat gaacaaaatt tagccaatat tatcagataa atcaaataact taccgtagaa 60
ttactgaaaa tagaatctac atcatcaaga ctaatattag cctgttcaaa tacaggaggc 120
ttatattccct tcttgaacca gtcattctca atgacctcag taaatgtaat ccgctgtgtc 180
atattattaa aagaaaaaca gatgccggtt gttaataata atatattgg tgaaatggag 240
ggaaaaagtaa aatattctgt atacactaga ttagactaga gtactatata attaaacaaa 300
aaaaac 306

<210> 669
<211> 363
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 669

gagagtcatc ttgatatatgac agctntggaa gtccttac gagactatgc tctagcatga 60
ccaagcttct tatgccatac ccagtatgc tctntgattt anagtaagca tgagaacctt 120
tgactggaca gatcaccaag tttaatatta taaaaggattc cttgtctttt agcaaagaag 180
agtgaagagt tgtccttngt ctgaacaata cacatcctt ttaaagggtga cattgtatcc 240
actatcacat aatcgactta ttctcaacag aatatgcttc aatcctttaa cangtaanac 300
attatctata taaggtagg gaggaacaca tactttacct acacanngta tcataccctt 360
ctg 363

<210> 670
<211> 505
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 670

atatgaatcc tgcangcagt natcccttag aatgcngccg gtcgnancct nggaccgatt 60
catgcagaca agggacncct gtanagcata catagtttt tgtcagctga actaatgagg 120
cgaacatgct catcatgctg acgcaactca actcatgggg ctccctcgaga taatgataat 180
taagcagaaa cctaagatgg cctgtcaatc ctatacagca ataattacag aacccctcct 240
tgccttgccc accctttggc taaccatgag taacacacga cgtcaacact aacgtatgct 300
gtgcccacac ctgaacaact ctactaatca ccacactacc tccttggaa ttgcatcatg 360
aggcttctca tgagcctcat cagggaaatg attgatatcg cccattattc ttgacaatag 420
gtgccagtgtn gcattatcta cgtgtccccca ctctcaatca ttcnacgctg agggcgttct 480
gctcgaccca ctaacctntc taccn 505

<210> 671
<211> 228
<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 671

cttattatta ttccaataac tntgtgtcg gaacaataaa aataatactc ataaacatat 60
tananagcat ttaacaatga ggaanaaaat gtcataacc atanaagaaa aaaagccaca 120
agaaaataaa aataaaatag tatntattnt ctnatctac ctncttatta cctaattagc 180
tcaatcttgc aaatttaaaa tgcacaattg accaaaaaat aacttgat 228

<210> 672

<211> 226

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 672

agcttgagat gaggaagtgt tgaagggtga atttcctgc ttttattgtt gaccacagag 60
tggtacctgn agatatgtcg cggnngtcag gagaccttgg ggacgtcagg tgggtgcta 120
ttgccccaaa ccaagcttga ccaatcccga cccaaaccccg gcatagtcgg tcagtgagaa 180
catgtgacgt acctaaggcag gcgagctct tgcagtcaca gataaa 226

<210> 673

<211> 296

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 673

caggcaacta actcctcttt canaaccatg ctatgtgctc gcgactggtc cctttttcc 60
tttcgcaact ttagttcaact attgctaccn catagagctc cgcgaaattt gttccggcca 120
tactcttcct tgcgagccct cttggctct tttcaaggg ctcttgcgg aattgcattc 180
tcttcccgta acccggcaca ctcctccga acgtgtgtag cggccaaactt gaacttcttc 240
ttggcaagtt ttgcctttcc taactcgcta ttgagagctt ggacttcttc gtgctc 296

<210> 674

<211> 113

<212> DNA

<213> Glycine max

<223> unsure at all n locations
<400> 674

gctgcaagct tggattgatt cggctcgaca agggattaat gnttagtaat ttaggctaca 60
acattgaaca aaagaggcaa atatgatcat catgcttga taaaaaaaaa atg 113

<210> 675
<211> 320
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 675

atgagaatgg ggttagatttg gagcatactc tcatactaaa caagtctata acattaatct 60
aaactcgctc aaactggttt tacaacgaan aatctaccga atcaataatt gactcctcaa 120
cacccaaattt accctagaaaa tggctcttgc cttcaactttg gtcactcatn ttccctcattt 180
gctcagccca agctgtccca taagtccaa atgacgattc anactaggat taactcactg 240
taatattcaa ttaccactaa atccagaatt agctnttcag acccctcaag cattacactg 300
tgtcaactcat atcactacat 320

<210> 676
<211> 348
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 676

aaaaaaaaataa tttaatcgaa ctgatcgaaa ttccggAACat cactctaATC tgggtcaaat 60
acaanaaaata aaaaaagaac accatatatt taaaaaaaaa aaaaaatttc tccacgggac 120
ctatgcaaca ggatagcaaa aaattcagtg cgctgagtga ataaatgctt cccaaaggac 180
gtaaatcgag gggAACGAAA caaaaatgat tcataatctga aaaaaaaaaaca gacaccaat 240
accaatgaaa gctttgcaaa aaaAGCATAG atcagcgcAT atcaaACATA gagcatgatt 300
acataatcag atgaaaacgc ggatttatata tacaatagta atatgagt 348

<210> 677
<211> 312
<212> DNA

<213> Glycine max

<400> 677

ctcctcatac agtggaaactg gcacaagtga ccagacatct gatacggtag agggtgattt 60
attattcatg tttcatagtt gtatattata atccgcacgta tatctataag aatgtatgtg 120
ggtagcaatt catcaagtga aacctctatt cttaacacat gaggaagttc acagtacaag 180
tgaaccattt ggtattctca atgacaataa gttaaaaata tcagcattca atgttagagag 240
ctcatccgaa gaagtccggg agatcttcat gggtgcatga gctacagatt ctgcaagagt 300
tattgaagtc ac 312

<210> 678

<211> 381

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 678

cgctctgnca ttcagnanag aaggagggtt gagtcccggt ttatggtttgc atcccacatc 60
aggaacgtta gcaatatcag tacttggtct tccgtatgag tttgaggagt caactgaggg 120
tatagtctca atggccagct tggttatagt atcttcttgc gtacctgatt ctggagagcc 180
ctgcttctgc ataaaattgt cgacagtatt atttctaagt tctggggca tggattcacg 240
ctcaccagat acagggattt ggttgatttc aataactgag tgctttgtt catcagcact 300
caacttattc agctgataga cagaaggcat agactgctgt ccagatggga gcctcttctc 360
agtgtttcaa ctgatggact g 381

<210> 679

<211> 381

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 679

ctaaacctaa atctcgcat actaataat ggtaataact gtattattct ggttttacat 60
ttntgttcta tgtggcttcc cagattatct cttcacctag taaataagag actaatctac 120
caagcatgat ctggtggtga naaagatcaa ccctctctag tctcaatggt acctttctca 180

caaacaaaagt tcagaggatt gaatcanaaa tcacattcac atgctcaagg gatgtgagcc 240
gctgtcactc acatcaatgt atggtatgtc tgtaatacgt tggtagttt atgttattgg 300
caaaatggat tgttcacaac ataaccacta tcataatggca cataactcac ctattgagag 360
catgatcaac accttcagta t 381

<210> 680
<211> 317
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 680

agctngaccg atcccgaccc aacccgcgaa nagttcattc gtgagttatga aaagaggcac 60
atgtaatcat cctgctcgaa caaatgacaa cactggggca aataaagagg gtgagaatgt 120
agaacaaacc catgctgcga ctgccattac tatacggaca aggttcccac caacccaaca 180
atgtcattgc tcaaccaata acaacccttc tccctaccta ccacccaggt aatcacaaag 240
gccatcccta aatcaaccac aaaaccatc ttccacacaa ccaatgctaa gcaccacctt 300
tagcacaaac caaaaca 317

<210> 681
<211> 151
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 681

cttgcacttg aacgccaatt gttatcatca tatactaattt gatcacatcc agacagctt 60
tccgaagggtg ttcttaattgt gtcggaaatn ttgcacagac tacaaaactc tggcgattaa 120
tatttctgac gaccattatt ctaacggatt t 151

<210> 682
<211> 334
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 682

acatgataca tgtcttggtt tggctggct caaagataaa ggagatgccc cacattttt 60

ccatgacaca natgcanaga tcatgatttg gaaacttcat gcaaaaactgg tcatgcatgc 120
acctatgtgg acactcaagt gtcaaacttt tatggtcatg tcatgctagg gctcangatt 180
tagatcaacc caatgttgcc aaaatatgtt ctttatcca tttgtgcatt catccgagtc 240
catttcgggc gttcggtgaa atttcacagt gttcaccctt caggtgtaga cacatnttt 300
ttcaaaaact agttatgatc aatgaatttt ttcc 334

<210> 683
<211> 407
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 683

gacaactctg atatcaagag acttacacac acacactntn tcctagtcga tcactcacat 60
anatntccat tctcncccct tggtttgag tttatgcntt catttgaat tagttaatta 120
cttatgtgag ttcttgattt attccctata tctctcccc tttggcatca acaaaaagcc 180
aaagtgtgta acaagtataa gacacacata tactattaat cattcacaag gcatacattg 240
aagaatataa accaatcatg aagcacgaaa catgaataga tcanatatat aataaccaca 300
tagtcatata atataattca taattgttca ttcacaccat gccaatatacg aaaatactaa 360
atatccaaat gtcataataa tatatggtat ttggataagt cactaca 407

<210> 684
<211> 278
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 684

agttccttc anaaattacc atatttgcgt ggtgactact aagagatagg ctaccaacaa 60
gatcaaattt gaggagaaag caaattgagg tcgaagatcc actgtgtcca ttctgcattc 120
agctgaggag agcgcttgcc aactatttt tcagttggag aaggatactc cgctatggtg 180
ggaatcatta tcatgggtga angtagtggg ggcttgcata aatcatacaa ggcaacactt 240
ncttcaacac atatatggag cgacagatgg aatgagag 278

| | | | | | | |
|------------|---------------------------|------------|-------------|------------|------------|-----|
| <210> | 685 | | | | | |
| <211> | 454 | | | | | |
| <212> | DNA | | | | | |
| <213> | Glycine max | | | | | |
| | | | | | | |
| <400> | 685 | | | | | |
| | | | | | | |
| cagaggtggt | ggttgttta | tgttatcgat | tttcactctc | tattcatatg | tttccgtatg | 60 |
| taattatgaa | aacttcactt | aatttatgct | tagtttgat | tcttattttt | ggctgtgttg | 120 |
| tttatttggg | gactggtgtc | cattaatatg | ctttggcag | tgcttttca | tttctaacac | 180 |
| gttcttgtgt | tctgcaatct | ttttgcatgt | aatttcttat | atacttagta | gttggaatgg | 240 |
| gcactgtgtt | atctttcag | gtctaacacg | ttctttgtt | ctgcaatatt | attgcacatg | 300 |
| atttcttaaa | taccgagtag | ttggtagata | ttgtttctt | taacagcata | catactgcc | 360 |
| tgtgccatag | agatataaaaa | ctgcttgact | cctgggtgtgc | acaatgcgat | tataactcaa | 420 |
| ctcactgtgc | ttgaatctgc | ggacttctga | aacg | | | 454 |
| | | | | | | |
| <210> | 686 | | | | | |
| <211> | 304 | | | | | |
| <212> | DNA | | | | | |
| <213> | Glycine max | | | | | |
| | | | | | | |
| <223> | unsure at all n locations | | | | | |
| <400> | 686 | | | | | |
| | | | | | | |
| agctntgctg | atttagttnt | cacttacgaa | aggtaaaag | tgagtctgaa | aagaggcaaa | 60 |
| ttaatcatc | ctgcttgaac | aatgagaaa | actggggcaa | ataaagaggg | tgagaatgta | 120 |
| gaagaaaccc | atgctgcgac | tgccattcct | atacgncaa | gttcccacc | aacccaacaa | 180 |
| tgtcattgct | caaccaataa | caacccttct | tcttacctac | cacccagtta | tccacanagg | 240 |
| gccatcctaa | atcaaccaca | aaaccatct | tccacacaac | ccatgctaag | caccacctt | 300 |
| agca | | | | | | 304 |
| | | | | | | |
| <210> | 687 | | | | | |
| <211> | 314 | | | | | |
| <212> | DNA | | | | | |
| <213> | Glycine max | | | | | |
| | | | | | | |
| <223> | unsure at all n locations | | | | | |
| <400> | 687 | | | | | |
| | | | | | | |
| gcttctctgg | gacctangca | naccttcaac | tcatccttca | cgatcaaagt | gtctactcgt | 60 |

cattggtcgc tntcctccct ccggagactta agctcgctat ctcagttcgg cattntcctt 120
ttggatctta agagttgctg attngaacct ttatttgac cgttgggctt gctcgagtcc 180
tgccctaagg gactacacct cttcatcttc ctccggtgcc tcaacttcct cccctttgc 240
gtgtgggatt tcagccactt acggtagcct ccaatgggcc cgttgtctnt gtctttctt 300
gcattatttc ccat 314

<210> 688
<211> 120
<212> DNA
<213> Glycine max

<400> 688

agcttgacca atcccgaccc aaccgggca tattcggtca gtgagaacat gtgacgtacc 60
taaggcaggcg agtcctggc agtcaacaga taaaaggaaa acacgaccac agagcaggga 120

<210> 689
<211> 224
<212> DNA
<213> Glycine max

<400> 689

cttcacccccc ttgtcttcaa cgtgaactat gaccattgtt ctatctccc gcgatgcttc 60
ttttcatgtc cgccgtgatg ggcttatagc ctaaaccata ctctccacga ttgtcttg 120
tatttatcag gctagttatg ccgacgttgt atttgctata cccatcctgg gttcataacc 180
gttccccaaac ataactcgaa ccatcattac cgctgcacatcg gaca 224

<210> 690
<211> 209
<212> DNA
<213> Glycine max

<400> 690

tctgtgtgggt ggtcgccaga ggagcataaa ccacacagtc tggcgacagg tgcagatatt 60
tgagtcatgg ccagttgggt taccaggta accaaggcat ctatgttacc ttcaagcttc 120
ttagtctcac ctgatgaaga tgaattcatg gctacttcaa gcactcctct aatgacaata 180
acatcatttc tggcactgaa ttgctggga 209

<210> 691
<211> 386
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 691

accaagtggc ctagaatttg atcctggctg tcgacactct ttaagtgaag agccgcacgg 60
ttggggagcc tatcctgtgc cagcatcaa gcttaaggg ctgttatgta agggagcggg 120
ttacgcatgt aatataaaac cattggtgtg ggatttagcct acaactaatg ctattctagt 180
tctctagata ggtgggtgac aggcatggaa gacggcgaaa aagttaacta taataccgca 240
tcataactaat gaaaacttcg catgctcaat gaatgcttaa taagatatgt ggctgcacaa 300
gaaaagtgac acttacgang accgcgtcta tagatcgact cttgccggat ccttgcattg 360
cgtcttata acgcatacat tctgac 386

<210> 692
<211> 187
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 692

gcttatattg taattgtta agttgtaaga ctctcgaaac accttgtana tcctgagnaa 60
aaaaagatta ngtgcttaat ttgtatatct gtctataaga cattaaggct agtttatgtg 120
catacaaaca tcaacaactc tacntaatttg ttagagccag aaatggctta atagtcaaag 180
aatactt 187

<210> 693
<211> 614
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 693

tcgacacttc anttctccat tgtatactcn cganatcata attcntntac tggctactga 60
cacncaaanc annaccggaa gaganaattg atgcgtctcg tacncgagat cctctaagtc 120

tacctgccgc atgccagctt gtggaagagg caaataacat attatgcgtg ttgctatata 180
actatctccc atctctaat caatccttg tatagtgtag gcttgtacac cacaacttg 240
catatgggaa gttttagggag gtgtacaact ctatcttac gctgtgagaa ttgttcatat 300
tggctgtggc tatccttgtt gaacatgacg gtttacgtgt tggcgatgg tgtgcgtgac 360
tcaccacatt tgctaaactt gtgaatgttc tctgtatgtga cgatcaagat tacgattgtt 420
gctgcataac gggttctgaa ggatgtatac tgtacgggtg ntgactctta tctactctat 480
tccacagcac gcagcaccca agctacttgt cctccggAAC cataaatcgt acgacgtatc 540
atacgttccg attgatcaat cgaatgtaac tatccacatc tcggttcgat catacctatc 600
accctaacta tacg 614

<210> 694
<211> 436
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 694

caagatgatg ggctcaacat atnaacaaac ccgctggcca atcatggagg ngggcggata 60
atgccataga tgtgagtagg ncgcaccggc ccatactttt gaaggatgga acgacacctca 120
gaaggtttat ctacgaagcc ctgtcaaaga cggatgatt tcccgtgcca gagaacatct 180
ttgctaatgc attccgttgt actgcgtgac atggaaacat gctcggcgga tagagatcta 240
ttgcacaaaa tgatacaaca cggctagcta gatgtcgat atgatggggaa ggaagaatca 300
catgttacat cggcttacc aatgatcaag tgctcataat cctaattcatt ggtttatcatt 360
accttagacac tattccaatg acccaacacc ttgcgtatg ggcagcaaca ttctattctt 420
acttatcgca caactc 436

<210> 695
<211> 425
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 695

agatgcgtgc ccatctatta aatgttgcga gtataatctg ctctattatt ttacaacacc 60

cttgtttaaa gtgttcggt aattatataat aatgaaatgg aaactttatg cttctatatt 120
tatgattcta tcgagttcaa ctgactataat accaatattt ttgtggata aatgcactct 180
aaccatctta attattacta aaaaaatgac atctaagaca gattgattag aatatgttcc 240
atcaaattat ttgtaaaaaaaa atatatagtt taaaacgatt cttagaaaaa ttatttcaaa 300
aaaattatTT ttgtaaaatg gtttttaaga aaatcgatt agaatcttta aatctgttaa 360
cttttttgg tttaaaaaaag acattataaa acnggttctc taaaaatcgt cttagaaaaat 420
ctatt 425

<210> 696
<211> 532
<212> DNA
<213> Glycine max

<400> 696

aacattcatc tccatgtact attagttag caacatgata ataatccacc cagcacgtac 60
atgatgcttc gtctcggttc ttatatcacc tgacgcatca agcttgagcg aaaagtgtga 120
agatcacact tcctactttt attcgcgacc acgagttggac ctggagatat gtcgcccggg 180
tcaagagatc ttggggaccg caggtggggt gctatcttcc aaaccaagct tgaccagtcc 240
cgacccaacc cgggcatagt ctgtcagtga gaacctgtga cgtacctaacc caggcgagct 300
cctgcactca accaataaaag aataatgacc caatgcagga cgctgcgtgg tgctggctac 360
tatggtcttg gtgatactgg atatggctta gctaagatac ctcggatgtat atcgatacaa 420
gctaaaatga aaccgcagct atattccttg gtatccaacc acggtggtat gttcacccctc 480
tataatggat ccgagttgca tctcttgcc tcttctactg gacggcgtta tc 532

<210> 697
<211> 455
<212> DNA
<213> Glycine max

<400> 697

ttataccatg gatgaataac agcggctcag ctacatcaaa aactaccaaa agaaaacttag 60
agttgacaag tattgcagct tacaaagttc attggatact ggaacaaaca aaggctcgac 120
taaaggaaaa agagtcattt taccttcaac ctttgggg agcccatgtt acatggatca 180

actttatgg gatggtatgg caatatgtgg tcatgttggt ttcctaaatc ttttataac 240
tctaacaatgt tatccaaatt gtcctaaaat tcgttagatta ctttcacctt tgaatcttaa 300
accaacagac aggccagaca ttgtctcatg aatttcaga ttgaaatatg aacaaatgct 360
ttctgactta ccaaaggcatc agctgctcgaaaagttgtt gttgcgcgta agtttagaat 420
gatcttgct gttgaacgta gaaatcaatt gatca 455

<210> 698
<211> 431
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 698

gcttgtaaga attgcaagat catcttcatt gactacagtt ngaaaaagat cgccaccaat 60
ataaaagagat gacaatttag agagtgtatcc aagactttca aatggatctc cactgaattt 120
attagtagac agatcaagat atcttaatga tgaatgctt ccacacgatc taagaagagc 180
accacccccc gtgttgtgg aagaatctat ccgctcaata tttttaatg cacctatata 240
atctgtcaga tggcctgaaa gtcgtcaact ctgaacttga agtcttgtga gtccatggga 300
aatacattga gcaagaatct ctaacagttc attaacctgc tcggttgagt ttgagattcg 360
agatataaat gtccttaag tcgcagagat tacccaaaga agttgaaatg ttttcttcat 420
gtcgataacct g 431

<210> 699
<211> 381
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 699

catcangaaa caatntcaact ttaaaagtgg gtcccaattt gattcctaag tttcaactta 60
cctttttggaa agtgacatca tggcagtttag gtcccgatctt tccatcgatgg attcagtcac 120
aaaacaaact tcaatatgtt ggactgtcta acacggngat ttttagattctt attcccaactt 180
ggttctggga accacactct caggtttgtt atttaaacct ctctcataat catatccatg 240
gtgagcttgtt gactacaata aaaaatccaa tatctatcca aactgttgat ctaagcacac 300

atcacttatg tggtaaatac cctatctatc anatgatgtg tatgggttaa acctttcgac 360
caattcattc tctgaatcca t 381

<210> 700
<211> 389
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 700

agctntgagc canaatcatg actcaccata taccttgacc cagggtgaga atgccaatcc 60
ttaccctcg aagcaaaaaa aagaagagaa cgaaaatttc caatcaaagg aaaaaggaga 120
aggaaaattt ccaatcaaag aggaagcaaa aaaggagaga aggaacattt ccaatcaaag 180
gaaaaagaga gaaaaggaaa ttctcaatca aagagtgcga gacagcaaaa agaaaagaaa 240
gataattccc aatcaaagaa tggagatag aataaaagag agaagtataa aagaagactg 300
ctcctggtca aagaanacag aagaaatgtg ccgagaggtc cttggaccag acgatatctg 360
aacaatacag aattgtcacc aaatgaaca 389

<210> 701
<211> 346
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 701

aacacaccaa aagattatga tgatgcattg ctcaaattct caccacttc aaattgagct 60
ttcaaaacta tcattgacatg tagaagaaaa ataaggattt caaatcagaa aatgtcaaga 120
gactttatt ttccagaacaa ttaccattt ctgtacata tcctataatt caaagaanaa 180
tatgcaaaatgt tacatgca aacagaattt gacctaaata tttaactaga aacccaaacat 240
acttaaaaac aaaactaaca aaactaacaa aacttaggaat accaaactaa cttaaaaaat 300
tactaaacca aaaccaaaga acaagtcccc catacttaaa caacac 346

<210> 702
<211> 416
<212> DNA
<213> Glycine max

<400> 702

agcttggcat caaagcgctc tattcagcat ttgcactcca cccttcgtgg ggatcttagc 60
agggttttt taccaatacc ttctctctcc aatatttgct tgaaatacat tacttaaga 120
attaatttct aaattacacc atttctata gcaaagtgg gttctgacat cccaaacttga 180
tatttttta tcattctaca cattctctct ccattgtttt cttacaatac actactttat 240
atattaattc tcacactaga tcacttcaa caaattcag aagatcgggg tcggacttcc 300
cttttttatt tataaaaacac tctatatatt attaaaatta aatattatat tatataaaaat 360
tatttctaatt taatataaaaaaa ttttagctatc tattaaatta atttatggaa tattat 416

<210> 703

<211> 434

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 703

tgataagtnt atgggctaaa ctgggttga ttaggggtga ttcataatta gtcataaacc 60
aattaacttg aanagttaaa tgacttataa gagtaatttg atgaaaatac ttttttgag 120
ccacaagatt atacgaaaga tttgataata ttaaattcga gcagaatatg cttcctcgag 180
ataaaaataa agaagcaaata gacttgagac aaattgcttt tggttataaa atgaatgagc 240
agaattttga atctctata gcggtaaaaa aatttagcag atgagttcan aatttttaat 300
acaataactt taacacctct tgattggaa aaaaaaatct ttgggttattt atctctaana 360
tccaagctcg taagttata aaaaaagata aatatacgatc attgaactat gctgatttag 420
gagatgacgt atat 434

<210> 704

<211> 420

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 704

gcttcttctt ggttctctcc ccatngaaa ccatcatttt tcttgagcac ttcattgaga 60
ggtgctgccatgtgctaaa atccttcaca aatcgctat aaaaacttgc taagccatga 120

aaactcctca cctcgacac agacttagt gtaggcatt cttgaatgc cctaacctc 180
tgctgatcaa cttgcactcc ttttgaactc aacaacaaac caagaaacac aacatggta 240
gtacaaaaga tgcattttc aagattggca tacaattgtt ctttctaag cacagtcaag 300
acagattta aatgatcaat atgcaaatac agtgaagtgc tatagataag aatatcatca 360
aagtacacca caactacact ttctatgaac tctctcaaga tatggttcat taatctcatg 420

<210> 705
<211> 401
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 705

ntataagtgc gggctggga gactaatgtc aagtgttcgc gatatgtgaa gatgatgttc 60
caagtacttc ggatttggtc cgaccatgcc ctccctgattt ccagctggga aattggcgag 120
tggaggaacg ccccgccatt tacgcaacaa gcataatgt aaccttacg gttttaaaag 180
ctctatagtt gggcttaggc ttcaagattt tcattttgtt aaggctttgt gtctttgtt 240
tttaattta taatacaagg atctntcttc atctgttcct ggtctctacc cattctcatt 300
catttgcatg ttacttctt tntctaaaac ggcagattcg atgacgagtt ccccgaagta 360
ctaatacctg ggacccgtct atcaacttcg agcaagaaat g 401

<210> 706
<211> 421
<212> DNA
<213> Glycine max

<223> unsure at all n.locations
<400> 706

ctcctacant tcatctctag catgcattnt tctttcttc tttaccact cctcacgttt 60
ggtttttag ggaaaaacat cataactaaa cgcgccacaa ggcaccccta tcgcaccaga 120
tccaaatcta taacgatggg tgatcaagag gagacacagg aacagatgac agccgacatg 180
tcggctctga aagaacaaat ggcctacatg atggaggcca tggtaggtat gaggcagctc 240
atggagaaga acgcccac cgtgccccgt gtcagttcg gtcacccaaact 300
ctcttggAAC tgtgccccatc ctccctaaAGC gtgttaggACG ggaagggacc actgggcacg 360

atggcacccct taccttgcata caaccgagcg gctaccctta ggatngccgc caactactca 420

c 421

<210> 707
<211> 450
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 707

tcgtgatttt ctggcagcat tgtgataaaaa gtccttttag acaaagctct cataacaatt 60
tcaggtgcca gtaagttaga aatgcacgtac ttgatacagg aaatggattt ngaaatttgtt 120
catcaagaat ctatcataga ccttggacga caaagtcaat attggaaattt cgagtaagt 180
caagatgtat taaaatataa catggtaaaaa tggatattca actgctttta ttngataata 240
cctgtgcttc tttagaatgta tctagtatga tagacctgat cattattttt ntaatttttg 300
gttgcgtatgg aactgatattt cgtgaaggca taactctaga tttgagtaaaa ttaaccagg 360
atctatattt gagctccaat tccttggcaa aattgtctaa catgagattt ctanaatccaa 420
tgatttgc tacatgactt actgatttac 450

<210> 708
<211> 330
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 708

agcttnnccc agtccatgag ccgctgaagt tactcttcta tcgnggngca ggtttccatg 60
tcgtgtgatt ccccgagatg aaacaaacat tcctcacttg tgccttcgcc acgagagacc 120
atgcacgttg cctgcagcga ctggtagatg aaccgtctag acatagccac atcttctaat 180
ctctctgact ctgacgaccc attctttgtt atggcattta tgcttaactcc cccatgactg 240
gctagtgat tggtttaacg ttggggccct cttcttgaaa ggatagtctc tccacactta 300
ttatgtgttag caccttatac ttgaatggcc 330

<210> 709
<211> 381
<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 709

ttataagtgc gggctcgga gacgaangtc aagtggcgc gatatgtgaa gatgatacgt 60
gctcatgcaa caattgttag ccatggctat acgagacatc ttgccaaaca aagtttaggtt 120
agcgataact cgcatgtgct ntttcttcca tgctatatgt agcaaagtca ttgatcctat 180
caagtttgat gagttggaaa atgacgccgc aattatactg tgccagttgg agatgtattt 240
tccccctgct ntatttgaca tcatgattca ctgattgtg atctggacag agaaatcaa 300
tgggtggtc ctatttatct accgaggatg tacccggttg agcgatacat gaagatcgta 360
aaagggtata cgaagaatct a 381

<210> 710

<211> 464

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 710

agcttgagaa ggtggcaaaa agcttataga aactttaaca tgtatttact atggagtatg 60
aaaagagagg acttgaacag aggttaaggta taaaacacca gttctgtggt gaaagtaaaa 120
acgaaatatac tggctcacac ttgattcaat gaattgaata gctcaaggaa atataagcac 180
ccatgatggt tgtggcatgt acttcaacaa ttaattcaat atagaaacta taataattga 240
tacgaaataa aatgtgtgga aatattgaga ccatacttca natgagtaag taatatgcat 300
cttgagggttc caagtatata ttgatggctt cacagattca tcccttgaag taaaaatact 360
aacaaggcaat ttgaagatca actttccagg tatgtaaaac acagaattag gatcatctac 420
agtcttatat agaactgcat caaccttagtc cattgattta atta 464

<210> 711

<211> 502

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 711

gcacccttg aaccctgan nnccctganc ccctgttgc accttgctat tcgtgaccc 60

tggatactca gccttgtat cactgttatca ccagcatatc tttcagctct attgaaagcc 120
tatctgcttc aaagaaaaat cgactcatag tcagggcggtt taacaaaaat cttctggaaa 180
ggactcattt ttgaattcat cttaagggtg ttcagtacat gtatggngtg agacaggttt 240
ggaaaataca attgttccag accccaagcg agttaattac tgctcacagg gggtagagt 300
tatgctaaat gtgtcagcag atggtacccc acgcaatgca natataatgt ccactattc 360
cgatgaatac cagaagctga agtactctgt ctctcttgaa atattcaat ttactttgtg 420
tgtgaacaaa gagaaacagt ccacacagat ggaacttgaa agaccagatc tgtctatacg 480
aatatatgag gagaaangcc cg 502

<210> 712
<211> 474
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 712

gctttttgga gtagaaacat gggaccaact catttttattt caaaaatgtt gtatctagtc 60
aaggcctaag attccataca agtttcctag cgatttctaa ttatgtgggc cattaagtct 120
atcatatgct gacaatagcc gagaaaccca tgaatttctt cggggggcggg gtaagtgtct 180
gccatcgccct tggccttggc taacaatcgg ggaagttcct gactccgtt caaggtaaga 240
gcaaaccgat ccatccacat ggttgcctct tggtgaaaga gtcgatcacc cttcctctag 300
cctctttntc cgcgataact tgggcataact cgccgcgat cctatgctcg tggggccgtgg 360
ctagacctaa ctcttcttgg tacttggcga tgatagctag catgttggtc tccgtctcgc 420
atagacgctg agacaagctt cttttgacc ttgaacaggc aactaactcc tctt 474

<210> 713
<211> 427
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 713

gtgcgggtct gggagacgaa ggtcaagtgg tcgcgatatg tgaagatgat gttccaagaa 60
ctctggattt ggtccgacca tgccctcctg atttccagct gggaaattgg cgagtggagg 120

aacgccccgg catttacgca acgagcataa tgtaaacctt tacggttta aaagctctat 180
agttgggcct aggctttaga gtttcattn tgttaaggct ttgtgtctt tgaatttata 240
atacaaggat ctttcttcat ctgttcctag tctctaccca ttctcattca tttgcattgtt 300
tacttcttt ttctaaaacg gcagattcga tgacgagtcc cccgaaggta ctaatacctg 360
ngaccgcgtct atcaacttcg agcaagaaat gaaccanacg gaagatgaag gagatgagga 420
tgtggga 427

<210> 714
<211> 289
<212> DNA
<213> Glycine max

<400> 714

atttccccct ttttctttt aaatatacca ttataaaaaa gggaaaaact tacgttagatg 60
tattaaccta tttgtggaga tttaaaacat ctgggttaag agatccatct aagagcctaa 120
ataaaatctg gaagacagaa caagggaaat gttaaaagaa gaagaagtga atttgcattga 180
atagaacgcg tgagagaaga aaaagaaggg acctcaattt gcccattggc tgcagcaata 240
tgaagaggag agtgacaatc atagagaagg gtcagagttc aagagagct 289

<210> 715
<211> 383
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 715

tcttctgacc cccgcgacat attctcaagt accactcagt ggtcaactaa taaaacgtgg 60
aggactgact ctttcacact ttctcacacc gagcttattg ggatatgggg cacccgtcat 120
atgtggact aggtggcgat cgggcgatgg cgcanacaa atatcccatt tccacaagcc 180
caggcataag cccaccatcc ccangtgccc acctttaaaa ttatccatc accgggtccc 240
catcaacctc tccaagctt cacaatatct aaacaattca attccatttg tcattgaaact 300
acctttaaaa cagagtgaag gtagaaatct ttacacaaga ttcatcana ctccacatag 360
ttttccaac ccacatacct cag 383

<210> 716
<211> 69
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 716

agcttgaagg tggtaaccc accattttcc atattatata tactggnaac gtgtctacta 60
tcatggaca 69

<210> 717
<211> 240
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 717

agctntgtat aatgagaatt gtccttcaac aagcttgaa anttggcatc tgaagtctga 60
aagctctang cagataagtc tgcaaaaagct ggaagtggtg ctgaagtaaa agatgcaagg 120
atgccagcta ttggtgcan a ggaagg ggc gcatcagctg ctctgatctt ggtcttcctt 180
gcctctagaa aatataactgg ttggtcattc gcattccaac angttcttat gatataagct 240

<210> 718
<211> 461
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 718

agcttggttc aaccncgtaa tccaaggaat gggaaattctg attgccaata cttcaacaac 60
atctcatagg gatgaatgac tcgggcatac tttaagctt tgcacggaaa atgttaattat 120
gaaattgaga tgccccaga aacaccattt cctagttaac catgcattag gtaccatgtt 180
caattatttt gttttgttgt tgggtgtttt ttttttagaa atgggtttat gatcccaaca 240
tgggtggctc atgggtgccta acacatgcaa ctaagaatgt agtgtgaagt ttcacgcttc 300
ccctttttt gttttgtttt gtagaggaaa acgcaaggat gagcaaacat gaaaacaaat 360
ggtatgcaat tntgcagatc aaaaagttt tggaaacgcat atgcatgatg atgccatgac 420
tcatgcanaa tggaggctg gaatatgata acggacaaat g 461

| | | |
|---|---------------------------|--|
| <210> | 719 | |
| <211> | 398 | |
| <212> | DNA | |
| <213> | Glycine max | |
| <223> | unsure at all n locations | |
| <400> | 719 | |
| ttgaggaga atgttagata atatgtttt tcttaataa ctttcttaat atacatcgta 60 | | |
| gactatatta acacattcta agtaatcgat tacgattttt ttttattttt tattttatga 120 | | |
| tgaatttcata gctgatcaca cacactttt cactcatata atgagactaa aagaataaaa 180 | | |
| tatacgata cacgcaatat aaataatgga aaggaatata aatttactgt gagtcgacac 240 | | |
| tttcaattat ttttatgaaa tatataata aatattcatt atcctaatac aattatgaag 300 | | |
| tttttagacg aattttctcc ttttcttgc gagacttctt tcatagtcggg cgatggctct 360 | | |
| ccttcaga tgattattct tatttcanaa acttttat 398 | | |
| <210> | 720 | |
| <211> | 516 | |
| <212> | DNA | |
| <213> | Glycine max | |
| <223> | unsure at all n locations | |
| <400> | 720 | |
| ccggtnatat ntgtaaacc ctggagtttgc gcttaccat cgatnaccca canananaan 60 | | |
| aanaacacat ntgtatgaa acacgatgac acaatcacaa cgacttgtgt catgtattgt 120 | | |
| actacggaga tccttaacga gtgaatctct ctctaggaga cacgtatgga actgtatgaa 180 | | |
| tattacgcat acctactta gagcgaagct ctactgcttgc agaagacaaa ctagatattc 240 | | |
| acacaacccc atcttgatat gtttagctca cccgatgaca tattacatgt taagtgtgct 300 | | |
| gaacttctgt tgtgctgaga cacatttatt gactctctat gataccgtcc tgcgctatgc 360 | | |
| acaggttata gagacttac ggtgacctgt ctaacgaaat gccagttgat atcgtgtcat 420 | | |
| ctataatcgg gcttatgatt ggacacgggc tcgaaactaa cgtatggcga agacaacatg 480 | | |
| tggactatct tgggtttatc tcactgatct gggaaan 516 | | |
| <210> | 721 | |
| <211> | 391 | |

<212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 721

 atcttaagtc acctgcccga tgcaagcttg cctcanagag gtccaggaag gacatggcat 60
 ccgaaggaac tagttccgct ccggaggtatg acagtcaccg cttaggagc gctgtacacc 120
 agcagcgctt cgaggccatc aaggatggt cgtttctcca ggagcgacgc gtccagctca 180
 gggacgacga gtatactgat ttgcacgagg aaatagggcg ccagcggtgg gcatcactgt 240
 gtactccat gccaaagttt gatccagaaa tagtcattga gtttatgcc aatgcttggc 300
 caacagagga gggcgtgcgt gacatgagat cctngntaag gggtcagtgg atcccgtttg 360
 atgccgacgc tattcgccaa ctcctaagat a 391

 <210> 722
 <211> 289
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 722

 tgctatgtgc tcgcgagtgg cacctctt cccttcgcag cttagttca ctattgtac 60
 cccatagagc tccgcagaaga ttattccggc catactttc ctgcgagcc ctcttggact 120
 ctggtcaag ggctttgcg gtaattgcat tctttcccg taacccggca cactcattcc 180
 gaatgtgtgt agcggccaac ttgaacttct cttggcaag ntgccttt cctaactcgc 240
 ttttgagagc tcggacttct tcgacccattt ccgaggcttc aaactctct 289

 <210> 723
 <211> 296
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 723

 atgcaaaagg tacgtatagc aacgtgcctt ctattcggtc aagcagactt gcacttcaac 60
 ttatcatatt tttttttct taaacaacat ttttaccta aaataaaata aggactctgt 120
 aattntatcc tattttaaaa aaatattttt aaatctcata tatttaaaag agatgtgaa 180

gtcagagtaa ttgattatgg aatttaaaaa tccagtttat aaaaaaaaaatg tgattatccc 240
atcgaatttc taaaaaaaaaa catgacacat cagtgggtt aataattat tggat 296

<210> 724
<211> 125
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 724

gtatgtgnntt ggtatttatg ttaaatataa agnttaanna taanaatgag agtttgtatt 60
aatatttaat agtatgtaaag gngatgaaaa aaataaaatat ataaataaaaa tgaattnatt 120
attag 125

<210> 725
<211> 392
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 725

agtttcttga ggaaggcctct taatgaagct tcttgaggaa gctacatgag ctgccttggt 60
aaaaacgttg cccagccttc gataaccgtt ggatcttcgt gtaatttggt ttgcagcttc 120
acaagacaat tgtacacgat ctgcctgttg ggatcttga gaagatgtct ggagtgtgtg 180
tgaagcttcc gttcccgaga gaatttctca tttaagcatt tcagccttg cttcgtgta 240
gcttaagaat tccttctcct ttcttcttc canagtcatt tctaaccgcc caagcatttt 300
ctccatcacc cacaaccacc attagccatc acanaccgcc attgttctcc attgagaccc 360
acattgaaaag gaacccttca accgaagcgg aa 392

<210> 726
<211> 447
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 726

ggaaggtttg tacatgacca aatcttagt taatcgtctt tacctanaac agtctttgta 60
tttgttaag atgcatgaag atagatcact aggagaacaa ttggatttgc ttaataaaact 120

gattctagat cttgaaaata tcgatgtcac tatatgatga tgaggatcaa gctttgttat 180
tgggtgctc tttgcctaag ggtaactcta atttcaaaga gactntattg tttgaaagag 240
actttgttc tcttgatgaa gtgcaggctg ctctgaattc aaaggaattg aatgaaagaa 300
aggaaaataa gtccttaca agtggtaag ggctgacagc aagaggcaag accttcatga 360
caaatacgtaa atctgataag aagaagcana agccagaaaa ccagaagaat ggtgaaggaa 420
atgtcttcan aatcagaggt catcact 447

<210> 727
<211> 329
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 727

gatggtgaga ctgactctaa atttgatttc tgtggaccag tcttagtgct tttgaaattn 60
tatcaaata tacccattgg ttttttatta aattgcgtt gaattaaact agataaaattt 120
tgcaatgaat tctctattag agcccattgt gagatgaggt caaattaagg tccatttg 180
acatacaagg tggctgccct attgtcaaa ttgggttgg cagtgtgtgg gtctcttga 240
ttttgagtct ttataagggt cgaaatgtt ttttagctt cttaaatttag ctcataaattt 300
ctttctaacc atgtcttata gaaaaatta 329

<210> 728
<211> 429
<212> DNA
<213> Glycine max

<400> 728

agcttggtgg agttcaaaga gaatcgtaga agttcgccc taagatataa gcctacacgc 60
accgacatga agagaaacac cctataaagg agaggcagaa gtgtggcca ccagcaagga 120
ttgcaagtaa aaggaactcc cttatgtcac atcaacaaga gttttgtcag cgcacgctgg 180
atgtgtgagg ggtgggtgc catgatccat gatgaagtcc ctcaagagca atcaaactgg 240
gtgcggccat gccctcctat gttcgagttg gaaaattggc aaattatcaa acaaccacaa 300
atttttgtgg caaacataat gtaatttgtt aatccaaacc ctatagctga gcctcggtt 360

gtctttgct attagatata tataaaatat aatctggctt catttttct tgactttca 420

tccctattt 429

<210> 729

<211> 265

<212> DNA

<213> Glycine max

<400> 729

tcttgagaa aacctcccttg agaagctaga gcttagctac acttaccctt ctcatatacta 60

agctcacctc cttgagaagc ttcccttaaga agattcgtaa agaagctaga gcttagctac 120

acataacctct ctaatagcta agctcacctc cttgagatga gaagctagag cttagctaca 180

caccccttat aatagctaag ctcaccccca tgacaaaaaa catgaaaata aaaaaaaaaagt 240

ccttattaca aagacaactc aaaat 265

<210> 730

<211> 254

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 730

ggcacctcac ataacctgtt attctctngn ccaaggaaa gattggagt aaaactttgg 60

aggccacga aacagacgac aaattatttc taatgtataa taaatacttt tagtttgata 120

tatttttta aatgaacgag aaaaaaatga tagattaaca taaatggaat gttctaacac 180

cccagtgcgg ggaggcttcc cgctatacga aggtatgtgg gaggggtatt ngacacagac 240

ttacccttgc ctat 254

<210> 731

<211> 559

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 731

ctcgtaactt aactcggtt cgtactcgca actgngctcg tatttctatc ctatatgtta 60

tcncacacgc tgtgtatTTT gttgattctg ttcaacccg ngatcctata natagacctc 120

acgcatgcca gcttaaatac caccagcatc aaagatctat ggtctgttga cgaaacctct 180
ccaaatgcga tcttccgca agaattacgg aaagatctta taattgacct tagccgacgt 240
atccatgaag ccattgcgac actcaaccta ttatacgacc agccttgaa atgtttcat 300
tcggagactc tcatctaccc cacccatgga aatttgagaa acttaggtg cctctccggg 360
gaaaaaacat atttttttc cggggctccc tcttgacca attgcctgt ggcaaggttt 420
agaaaagggtt gaagattaaa acactcgga cggctggc cccccgggg actctaaaac 480
tggaaagggtt ggcaccaagg atgggcccc ttttggggg ccctctattt ttggggggcc 540
ctttcacccc cagcggtgc 559

<210> 732
<211> 329
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 732

gagaaaactt ncttgagaag cttcttgag aaaactttct tgagaagcta gagtttagct 60
acacacaccc ctctaataac taagctcacc tccttgagaa gcttccttga gaagattcct 120
aaagaagcta gagcttagct acacacaccc cctataatag ctaagctcac ccccatgcaa 180
aaatacatga aaatataaaa aaaaagtccc tattacaaag actactcaaa atgcccttga 240
atacaaggct taaaccctat actactagaa tgggccacat acaaggccca aaagaaggaa 300
aaccaattcc tacatttacc aagaagaat 329

<210> 733
<211> 470
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 733

acactcttag gatttgccata gtttacattt ctgcttaca ttcataatag cttatccct 60
ttaccttcca ttgtcaaacc gccttagatag cttccctttt accaattagt tntttacctt 120
atctttcaca cctcttttag tgtttatttgc tagnttca accatagttt ctttacctt 180
ttgttntcaa acctccaaca agaaagaacc acaacttagg aaccaatatg agtcatcatt 240

catctagtgg taatggcaag ggtactagtc ataaagaccc tttatctaga atcttagatg 300
agttgagttc cctcacgtta tggaaagaan aacaagagag aaaagaanaa ggaagaataa 360
gagtggaaaga aataaatcat gatgaaagaa agacaatatg agaggaagaa agaagaacaa 420
taatgaaaga aatgaanaga gaaaaacatg cctnctatag tagtcataac 470

<210> 734
<211> 429
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 734

atgatttggta ttatcctcta nggcaatcag tattcagtat tttaaattcc ctctcaaaaa 60
attgcaggct tggcggtt gccacaaagc tgccatatat tataagaggc ttagtagagg 120
tgcaatagtt acacaatgca gatggagggg gcgcatagcc aggaaagaac ttatgaaaact 180
gaaaatggta tggtttacca tgattcttat acattaagta gatccttcag aaatagatga 240
caaaagatgt aacacgtcca tcctacaaaa cttatggcac aaaaagcaaa agatactatt 300
tgatcaattt ttaattgtaa aattgagttt caaattatag atatatacaa acaatattca 360
tatTTTGT ttcatattt tcttcatgaa gggaaaaaca atagaagaac ttanaattct 420
cttctgatc 429

<210> 735
<211> 371
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 735

gcttattctt ncatggctta tttccttagag gatgggcct actctcacct ctccctcctt 60
atcttcgtt gtaactccat ggctgaanat caccattgaa ggacccatt aaagctcaaa 120
gatccaacct ttataaaagc ttctcaagaa agcttccatc aggactttg gacatcatag 180
tgcctacagc tgaggcatca aggatcatct tggttngagg tttcatacca ctatggaaga 240
tatgcatatg tgcactgtca tcanagttat ggttatgca cctctgcaac aaagctttaa 300
atctctccca tgtctcacag agaggtttgt gagatccttggataaaagtc atgatgaagg 360

ggtttacact g

371

<210> 736
<211> 335
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 736

gagaaagtgt ggaagagtca gtcttcctac ttttattcgt tgaccacaga gtggtaacctg 60
gagatatgtc gcggnggtca agagaccccttggacgtcaa gtgggggtt attgccaaa 120
accaagcttggaccatccc acccaaccca ggcatagtcgtcagtgaga acctgtgacg 180
tacctaaaca ggcgagctcc tggcagtcac ccgataaaag aacaaagacc acaaagcaag 240
gaggcttggatggctgg ccagctatgg atcttgagtg atatatgggt tatggcctct 300
ggtaatcgat tacaaagggt gtgtaatcga ttaca 335

<210> 737
<211> 504
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 737

atccttaagt cacctgcggc atgcaagctn gttcgacat cgntcgcgtg tatgacatcc 60
actccacaag gtttgaagta gaggagacct tcaatcctat tacgcaacgt ggcggacaaa 120
aatgggcagt taacttgaat ggtcattatt gtcaatgcgg aaggtattct ggccttcact 180
atccatgttc acatattatt gcagcttggat gttacgtgag cctgaactac taccaatata 240
tagatgttggat tatacataat gagcacatct tanaagctta ctccccacaa tggtggcctc 300
ttgggaatga agcggctatt cctccttcta atgacgcattt gacacttac catgacccaa 360
ctacaattcg tgcgataggt ccggcaaaat caacaaggat aatgaatgag atggattgga 420
tcgaaccatc tgaccaccga ctAAAATGCA gttagatgtgg agccgaaggc cataacangc 480
gtcgctgccc atgcaatctg agcg 504

<210> 738
<211> 411
<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 738

ttcacccgacg aaaggaccac agtaggtcta ataagagaca aatctgatca tcatgctttg 60
atacatgccaa aaaaaaacta cggcaaatacg agagggtgag aatgaggaag aagcccatgc 120
tgtgactgtc attcctatac agccaagttt cccaccaacc caacaatgtc attacttagc 180
caataacaaa cctttcctt acccaccgccc agttatccac aaaggccatc cttaaatcaa 240
ccataaaagtc tgtctaccgc acttcanatg acgaacacca cctttagcac ataccataaa 300
caccaaaacaa gaaatggaat ttgcagcgag aaagcctata gaattcaccc caattccagt 360
gtcctatgct gactngctcc catatctact tgataattca atggtagcca t 411

<210> 739

<211> 420

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 739

agctctcatc tagccaagaa tttacaaaag ttttacaaca taacctaacg atttctaatt 60
atatggccca ttaaatttat catgtgttga cagtaattga tttagccgtg aatttcctct 120
ggggctgaac acacttcggc catggccctt gctttggcta gtatcgccg gaggtcttga 180
cttccattta aggtcaaggc gaacctatcc atccacatgg tcgcttcttg atgcaatgca 240
tcaatcaccc tcctcttgc ttcccttctcg gcgtatgtt gtgcgaagtc ctcttctatc 300
tttgctcat gggcanaga ctgggttaac tcttctttgt actgtcctat tatanctagc 360
atgctctgct ccgtggcttc taagtgttgg gccaaacttt tcttggatct tgagcaagct 420

<210> 740

<211> 427

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 740

tgtatcgatt cgctcctagt caggatcaca tcatcttgct cctttcttn tcatttaact 60
tttggctttt gtttccttct ctcttataat atgtttggg gtataataat tcataatgt 120

atgtatgtgt cggaacctac cttcgtaa gagggcgagg cgaaaagcca aaggagcatc 180
ttccaaaaag gaaaacccgc gggagtgcacc accaacgttt actcttagaa aacatttagaa 240
aaaccaaaaa aaaaaggctcg aaggctcgca aattttgaaa atgagggttt gggagttgtt 300
tacacacgag gaaggtattt gcacccacg cactcgac aaggatggc aaccttaat 360
cgagtgtgca naacatgaac ttcaaaatgt gtatccc tttcatatnn gtttttttat 420
ttctttg 427

<210> 741
<211> 500
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 741

ttaagtccacc tgccgctgca agcttatgtat tttttanaaa ataattactt ctctgtgagt 60
ctccatgaac ctaataaata tgaggagaat gaagatttgt gagtctccat gaacctaaaa 120
aatctgagga gaatgaagat tggggatga attataaaat tgtgatcctg tctttataa 180
acctaactg acccgctntt ttcagaaaa aagataagaa tagttaaact ttgcccgtat 240
catacatatc actctaatan aaaataaagc attggagaga ccagaacata tgagatatga 300
atcgaattct tatcggtta tactttacca taaaagtcat attcaatatt atgtgaacca 360
ttgatntaga tggcttntt cttaataact catcattaa cagagttcag aataatataa 420
attgatgaat aaatcacatt cgcgcttctt atatactgaa cctattaatt caagccttaa 480
gctaataaa gcaacatttc 500

<210> 742
<211> 412
<212> DNA
<213> Glycine max

<400> 742

aatgtaataa gttaaacata aagctattga ttgccgtaat caattggcct aaaggccaag 60
tagagtatat taaatttcag gcaaaccctt attcgaatc aactggaagt cgatcagcaa 120
atctttaaa ttaacgaatt tgcctacttt caggaaggaa aaaaaaaaaa agcagtgtaa 180

tacctaata atatctgaat ttatcatacg tatttgctac aatttagattt tgtaaactgt 240
aaatctctct aatataattc ctttatgaaa gatgattgta acattaaaaa atatattttg 300
ggtcagcagc ttggatctt atagggggtg gtgccaacct agttggact actccctcga 360
ttgtatcagc tgtctcacca ttcatatta aaataaataa ccaggatcaa tt 412

<210> 743
<211> 361
<212> DNA
<213> Glycine max

<400> 743

gagactatgt ttggcaaaga aaaactatga agaatgcgtt gccatggaaa aactacaacg 60
cctactctct tc当地agaaa taaagaaatt tcaaaaattg aaaaaagata tgctgaagag 120
gtgaatgaat taaatgataa aatccaagag atagaagata aatattatgc aacagagaaa 180
tggtagcagc agagaaaaaa cactattaag aaattattgg agtacaagtg tgaggtgaag 240
tcccacatta aatagaagtg gaaaagttga gcaccatata agtgaggaga agacctataa 300
atctaagtct taaggtttg agttaaagtg tggattaa atcccttatac ttgttactca 360
t 361

<210> 744
<211> 386
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 744

tggtcaccat tntcaaggat agaattcata agactaagga aaaggattaa aaggaggtca 60
atgatgatga tcatgatgat gatgaagact atgtacctaa atatgaagag agacttggat 120
caaattctat aagtgaaaat taccagcagc acaaatatga tgaattttct actacaaatg 180
atcttgagtc acggaaaga aagataactc ttgaaaattt tcaatataat gctaataac 240
attattgtgt taataaatga gtgggttgtt tttgtggtta ttgtgaaaat tntcaatata 300
ataccatagg aaattgactt catntcttg atctatctac atctttctta anaaaaattc 360
aaattcactc ttttgaana aaatag 386

<210> 745
<211> 244
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 745

cataaaaaagt cttgtggaat tgattacaag gatnggtaa tcgactacca atgacaagg 60
ttgaataaaa atcacaagat gtaactcttc aaatggttt caggctattc tanagggtat 120
aactcttcca atgggttcca ttgaccacac ataaagagtc tataaaagcg cgacccttgag 180
tngcatattt agatctgagt acaaactttt acatctttt cacacaacct ttgaacatct 240
tctt 244

<210> 746
<211> 451
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 746

ggcttctaca atctccnct ntttgatgt gacaaccctg atatcaagaa acacatgcac 60
atacttttc cttagtcgatt actcacttaa ttctccatat tctaccctt tggttttgag 120
tttaagcttc acttggaaatt aagttaatta cttatgtgag ttcttgattt aatccctatt 180
tctctcccc tttggcatca acaaaaaacc aaagtgcgta acaaataaa aacatacaca 240
aataactaat catacacaag acattcattt aaaaatctaa accaatcatg aagcaagaaa 300
catgaataga tcaaataat aaaaaccaca tagtcatata acataattca tatttggta 360
gtcatactat gcaaataaaa gaaatactaa atgttcaaatt gtcataataa tatagccaaa 420
tacacggcta gaaatcaaag tactaataat a 451

<210> 747
<211> 476
<212> DNA
<213> Glycine max

<400> 747

cgcgcgcgcg ctcttattag agaacggtaa cgtcttgtac tattatacac acgaaaactat 60
atctacagga tgccacgtat gagtgattac ctcaataatc aggcgcttag actcgatctg 120

agagaggaga gacgtataaa ctcataact tctctaggat tgaagttctg atctatatcg 180
gaacattatac ttctgacgacg atacctgc acactacgct cagcataccca gacctctctg 240
cattgtatac gacggaggac ctccgagttac agttatcaa cataacctat ccaattgaag 300
cacatggact tttcaataaaa aagcacgaca attccattat tacacgtctg ctcattgaga 360
aagagatctc taggccgtag atatatactc tcataattcaa ctttagatgaa tgtagaaaaaa 420
atgtacttat agctgcataa tgatcatgtg ccggatctaa caatacacgaa gatgcc 476

<210> 748
<211> 337
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 748

tctccaccat tntcttataa atagggggag aagtgaagag gaatttcgtt cagccctcct 60
ggtaattcag aatcacttaa aattagtgaa naaaatttgtt tccgtgaaga anatccaagc 120
cgaggcgctt ccgtaacgtt tccgtgggtt atttcgcgaa ggtttcggc cgttttcgaa 180
cgctcttcat tcgttcttcg tcgntcttcg gtcttcaacc ggtaagttcc ctaaatcgaa 240
ctttcaatt cattctatgt acccttagtg gtcctcattt gctttacgt gcttcattt 300
acatttcctt tactttcgtt acccccgttt gacgtgc 337

<210> 749
<211> 489
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 749

agcttgcgtc agcggtttagg cgagacagag accatcatgt tagctatcat cgccaaagtac 60
gaagaagagt taggtcttagc cacggccac gagcatagaa tcgcggatga gtatgctcaa 120
gtatatgcgg aaaaagaggc tagaggaagg gtgatcgact cttaacacca agaggcaacc 180
atgtggatgg atcggtttgc tcttaccttg aacgggagtc aagaacttcc ccgattgtta 240
gccaaaggcca aggcgatggc agacacccatc tccgccccncg aagagattca tgggcttctc 300
agctattgtc agcatatgt agacttaatg gcccacataa ttagaaatcg ttaggaaact 360

tgtatggtct cttatacctt gactagatat gaattcccttc ttgaaatana atgagttgg 420
cccatgtttc tactccacan agcttgcata aatcanatca ctcctacatc tcatactctag 480
catgcattt 489

<210> 750
<211> 451
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 750

ggcgaccaggc tcgcccaggc gagtaagggt gcttcctcca gaagcaacaa ccttctggag 60
gaatcttcta gagggcccaa gtgggcctga ttgctatttg tacctccctt ttactaaat 120
ccatccccctt ctatttttt ggttaattctt tttccgtaac gttacgaaac ttacgaattt 180
tcgttaacgat acttattttc cttccgcaag gttacgaatc cttacggatt atgtatttac 240
tctnttttag ctntcgaaga agttacggaa acttacggat tgcgcanaaa caccttttt 300
cgatttccgc cacattacgg aattcacgg attgcgcaag cctgcttcct ttgatttt 360
gacaggcctc gggacttcat tcatttgca accaaggacg ccaagtatct cgaagcggcc 420
aatcaaagggt tgtatatacat caaataataa t 451

<210> 751
<211> 141
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 751

tactgaccat tgataatatac acaagtgagt ttattcagaa attagagttt atgtctttat 60
cttgtgagag tgattctcct aaattcttga gtgattanag aacaccctgc ctgtatcaaa 120
ggactttaac aacctttgag a 141

<210> 752
<211> 131
<212> DNA
<213> Glycine max

<223> unsure at all n locations

<400> 752

gattacctct ataatcaggc tcttagactc gatctgagag acgagagacg nataaactca 60
ttatcttctc aaggatagag tgtctgatat atatccaac aatatcatct gaggtcatcc 120
ctgcatatac t 131

<210> 753

<211> 427

<212> DNA

<213> Glycine max

<400> 753

tagggcagg tacgaacagg ttctatgccc atgatcaatc gatcaccacc cccgcgttcg 60
gctaaagata ttaaagaagc ttccttagga ggagcctag tatctctaac tttgctctt 120
aatttcctgt ttcatacttg ttcttttct tgaactatat cctgaattcg cctaagttt 180
tatgcaatta taggattta agagaaaaaa tataacaatg aataacacaa ttttgtaaag 240
gattttcttc accaaaaaaaaa taataattac ctcgcttggg cgagtggcca gctcgcttag 300
gcgagcatgg ctatggtaa aaacataaaa aggggagggg tgaagccatt ttcaccctat 360
tcttgccaa aatcaaaacc ttccccaga gcttacgggaa gccaccattt gcagcagccc 420
ccaagct 427

<210> 754

<211> 411

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 754

catcacatgt ggactatgt ggcggtcgaa cgatggtgca caacaagttt tccacatcca 60
caaatcgcc ataaacccac catcccctgg tgcccaccc caactgagct cacgtactcc 120
cacgtacccc atattctcgat ttctctcaac accgggtccc catcaatccct cccaaagcttc 180
cccaacattt acatccaaat catcacaaac taacaaacca agcaaaacag 240
ggcaaaggca gaataactctg cccaaaactc aaaccanaat cacagcttt tctcacttaa 300
agacccactt aacatttcct tcattccat tcgttaaccg gtggatcgac tcgaaaaattt 360
tactgaaatg ctctagtaca taagccata ttntgaccgg tggatctac t 411

<210> 755
<211> 268
<212> DNA
<213> Glycine max

<400> 755

catgccttg acggcaaccg cccacacgac acgtgagaga tcgacccccc tgtacagata 60
agccccata cctgtcatga tatttttcaa gatatggata ttaacctggc ttacagctgt 120
cttttggac gcccgtggat gcactcagtggat ggagatgatc cctctacact gcaccacaaa 180
ttgatagtcg tagtacacgt gcactggggc attgtgtctg tggacgaaga aatcttggta 240
agatgccat tctctatgcc atatgtgg 268

<210> 756
<211> 448
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 756

gcttgcataat aataatcaca gtatctgcat atttcaattt agtataatggaa catgtgttct 60
tccttaagag agggtgttat ccctgtgaa gatcctgcat actacctgca cccaaaggaa 120
cataacacag tcattaagaa tcagtgttct gagtataatca atgaattntg tgtccacaag 180
cctgtggcag aaccagtggc ttgtgtcaag ggaccaaggg attataattt ttcacgtatc 240
cagtcgtgct taaaagttgt ctcaaatgtt gatgaagagg ctgatgttga acctacatct 300
cctgaanagg aagggataga atgtgataat ccagaatctg aaagtaggtt agaatctgac 360
atcaagtgtt gttatggtt tgaaactgtat atataattgc atgacgttctt atagcagcag 420
aatctgaagt gaaaacttaat agcctaatt 448

<210> 757
<211> 454
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 757

agtttagggta gtgaatgata ttctgaaggc gtgttattgt gtaaccatt aataaagtgt 60

ctcgattntg ctataaaact ctgaaatgtg cttgctttaag attntatTTT 120
atTTTggTAC agaAGATTT atTTTTTGA aacaacaAGT attgacaACT ttCTGGCTGT 180
agtaaataga agaaggAAAG tggaAGAAAT aggAAAATGA taaACTgcAC ctCCtttAC 240
ttaatcttaA tgattctcAG ttgtttGCT ttccTCTTG caATTGTCTG aaACCAAGGT 300
ggaggtAGAA gTTAAATTc cataaAGCac ttaaAGAAAC tattTAATAT cCTTGCgTC 360
atTCatTTG gatGGAAAG ctatatacat ttAGCTAGAG catGCCATTA ctGcATAAAG 420
aataccCTAA ttaaggatAG attatgAGAC acAC 454

<210> 758
<211> 88
<212> DNA
<213> Glycine max

<400> 758

tgtattgAGC attcCTTTc tgtatCTTcC gTTTgCTTA gTTtagCCCT gtaattCTAA 60
tatAGATTAa gagAGCATTc agCTTgAC 88

<210> 759
<211> 172
<212> DNA
<213> Glycine max

<400> 759

tCTTACATAT tacCCcATTt atgtCTcaAG attTAGTgAT tcaAGCTTgg CCTCTTggTT 60
gagCTCTTAA catAGTAGAG gaaAGAAAC tCCCGAAATG atgGAAGAAg gATATGCTGT 120
gaacAAACC catATGTTCA agggAGCCAC ctataACTAT ggGAAGGAAa AA 172

<210> 760
<211> 315
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 760

gCTTCTTATC caggCAATTc ttggggNGA agCTCCTTCT tcCTTggCTT attCCCTAGN 60
ggatGGTgCC tCCCTATCC tCTTCTCCTT tgCCttCCGc tgCATCTCCA tgATGAAAAA 120

tcaccattga aggacctcat tgaagatcaa agatccagcc tccatagaag ctccacaaggc 180
aagcttccat caagttatga ccatttgaat ttctcgagat cttccgtgn tcaatttcgg 240
gcgtctccat atgtcatgtg cctgaatcgg acctccgtaa tataatttat gaccattcga 300
acttctctag agctt 315

<210> 761
<211> 410
<212> DNA
<213> Glycine max

<400> 761

gcagatctgg tcttcgccag tgaaaggatc aatgtgggtc cgaaaagagg caaatttgat 60
catcctacta tgacgactga gaaaactggg gcaaataaag agggtgagga tgagggagaa 120
acccatgctg tgactgccat tcctgtacgg ccaaatttcc caccaaccca acaatatctt 180
tactcagcca ataacaaact ttctccttac ccaccaccca gttatccaca aaggccatcc 240
ctaaatctac cacaaagtct gtctaccgca cttccaatga cgaacaccac cttagcaca 300
aaccaaaaac accaaccaag aagtgaattt tgcaagcgaga aagcctgttag aattcacccc 360
aattccagtg tcctatgctg acttgtccca tatctacttg ataattcaat 410

<210> 762
<211> 449
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 762

gcttaataac aatgcctaag gaggagcatt acatagcacc catctggtc ctgttaggga 60
taaacatgtc gtgaagtac attcaacatc tggttataaa ggactttac catggcatac 120
tacagtgact aacactgtat tcagaatcat tntcatccct tttatgaact tcccttttt 180
ttatgaactg cccttttt tagtcttgc ttccccaccc ttaaacatct gcattntccc 240
attttagttt cttgtttttg tgaggaatgg cgaaaattat gttttatatg gacaagactg 300
ttgggtggaa ttggaaatgag aatcaacatg gtgctcatac ttttgggtttc aaataaataa 360
aagtggccat gctactaata ttatnat tatggtgctt ctattaataa agtacacctt 420
gttagaaaaac caaaatatacg aacagaaaaat 449

<210> 763
<211> 287
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 763

tcttatccaa cgctcatctt ggtggtaag ctccttcttc catggcttat tccttaatgg 60
atggcgccctc ct当地cacctc tttcccttg tcttccgcta catctccatg gtggaaaatc 120
accattaaag gaccccattg aagctcanag atccagcctc catagaagcc ccacaagcaa 180
gtttccatca gaatgtccac gtcttagag ggctacacgc ccatgccttc agaggactac 240
acgcccctcgc cttatgagga ctacacatcc tcaccttag aggacta 287

<210> 764
<211> 435
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 764

ctgtgagaac ctgtgacgta cctaaacagg cgagctcctg gcagtcaacc aataaaagaa 60
caaagtccac gaagcaagga ggcttgtgtg gcccgtggcc agctatgtat cttgggtgg 120
atctggaaat tagcctctgg taatcgatta ccattcatgg ataatcgatt acaggggtta 180
aaaatggaga caggatggta aatggcctct ggtaatcgat taccaaggga gtgtaatcga 240
ttacacaggg tgataggca ctggtaatcg attaccagct ggggtgtatc cattacacag 300
ggtgataggg cactagtaat cgattaccac ttatgtgtaa tcgattacac agtgtatccc 360
ttaattttca atgtgcanag gctgtgtat tcgttttgg caccggtaat cattacatac 420
tttggtatcg atacc 435

<210> 765
<211> 318
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 765

tcatgatgat gaatgaagtt gattcaagta gttntgatga tatctaagat gatgacaaan 60
agcccaagag aatgatttca agattgagtt aacaagattt atgaatcaag agaagttga 120
tttcaagatt caagagaaga tgaattcaag attcaagaca agaaatcaag aagacttcat 180
aagggaagta ttgaaaagat ttttcaaaaa acaaacatag cacaatttg ttttcaaaa 240
gagctttct cagaattgtc taagttacca gagttttac tctctggtga tcgattacca 300
attaactggt atcgatta 318

<210> 766
<211> 395
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 766

agcttctaac ctttccttc ctttctacca catatgtgga gttattccac atacataaaa 60
ggccaccagc agcttccaca gatggaacaa aatcccaatg accagtggag tctcccgaaa 120
tggcctggca aatacttttta ttaaagntct ccctcttggt ttcttgagg cagacaagat 180
gcactttgtg cttacaatga gccttctaac agcagcccac ttgactcccc tcccctaacc 240
tctagaatta taggagagaa ttatcataat tgctgagatt taattccctt ttctgttgcc 300
atcaaatcat ctttattctc catatgcagc agtagccct taaccttgct atcttcttnc 360
ttataagaca agcccatttc cttcaagatg tcaca 395

<210> 767
<211> 432
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 767

agctntggag tttccaagtg ccaattcgtc ttcttctta gtccagtctt ctctggctt 60
caatccatca gtgggcttcc ttctgtgtc cagcatcttggatgttccc agcctttgat 120
gacagcttcc caggttctgc tatccagtga tttgaggaag gccaccatcc ttgctttcca 180
gtattcatag ntggttccat ccagaattgg tggtctgttc actggccttc ctctttctc 240
catgttcatc agaatttatac tcccttaggtc tcactcagtg atttcgagtg cctgctctga 300

taccaattga aattctgata ccaatgccag atgtcgtaa ggtatgtcacg acatcacgct 360
tcagaacatg cagattatct ctgagtggat gaacacgata aacaagtata taacacaaga 420
gaattgttta cc 432

<210> 768
<211> 429
<212> DNA
<213> Glycine max

<400> 768

tgtctcagcg tctatgcgag acagagacca acatgttagc tatcatcgcc aagtaccaag 60
aagagttggg tctagccacg gcccacgagc atagaatcgc ggatgagtat gcccaagtat 120
atgcggaaaa agaggctaga ggaagggtga tcgactctt acaccaagag gcaaccatgt 180
ggatggatcg gtttgctttt accttgaacg ggagtcaaga acttccccga ttgttggcca 240
aggccaaggc gatggcagac acctactccg cccccaaga gattcatggg cttctcggt 300
attgtcagca tatgatagac ttaatggccc acataattag aaatcgtag gaaacttgta 360
tggtctctca gaccttgact agatatgatt tcttttttg aaatgaaatg agttggtccc 420
aggtttcta 429

<210> 769
<211> 466
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 769

gtcacctgcg gcatgcaagc tngagaatat caatgcgtca nagtcgctat ctcaatcacc 60
tttggatggatg tatcaactca caatggagat gggaggagat ctccatgacc 120
acatcaacaa gttcaatcgg ctagtaagtt aactgttgaa tgtggatgt aaattctcta 180
atgaggagca agcgctttg ttgttggatc cactacaaa gtctccata gctttggatc 240
aaacgttgct tgtggaaaga tcaactttga atttggatga ggtgactgtc gctcttagag 300
aanatgatga gaattgaaaa tgctgatgtat gaacacaatg caatagctgt gatggaatct 360
gagcgagggaa ggaatcattc aaggagacat gatggtctaa gaggaagatc acaatcgcaa 420
tcgcacatccac aacgagatat gagtaacatt cactgcttctt attgtg 466

<210> 770
 <211> 345
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 770

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ctgaggatag agacttccta agctatatat cttctcttc anataagttc tctaacttcc 60
tagctatctc actctaagaa gtggattcac tcttgccttg gatggtagg aatgaaggct 120
cctaccctta ttatactac tccacccca caatgaatgg tggagattac ttgtatccta 180
gggtggagat taattctcta gaattctcca cacattctag gagtctctac acctttctac 240
tctctttcat atcattccat aaggttccag aaggttccac acatctccaa aatatttcag 300
agggttccac attcttccac aagcttctag agagttctac actac 345
  
```

<210> 771
 <211> 465
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 771

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agctntgcaa ctcagacacc tcaacaggag agatccgacg aggctgttgc tcctctttag 60
cctacacctg cacaggttga accagtgcac gctgatccac attctccagt ggcagatcca 120
tcttctcccg aacttgaagc agctccccca tcttcaccta ttattatcat ctctgaagac 180
cctacagagt caacatttg agaagctgtt gctctctctg attccctgt ntntcatctg 240
atgaatgagg aggagacaca ggatcagtca caggattctt anattcctgt cttctgttt 300
atgttgacaa ttatcataac tattatattn tagtacattt ttttagtgaa ttttgggtat 360
ggttatataat acttgtgttt tttggggaaa gtacgatgca tggtttgaag catacaggaa 420
tagttaactt gatcttgatg aatatgtatg gaaacacttt tatca 465
  
```

<210> 772
 <211> 397
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations

<400> 772

tcattgcagt cttggaagcc agattagttc atggatgaga tccccactt tggctcaa 60
ccttgccctg ttggaaatgga tctgggttgg tttgggttg gggttctcaa ttgaatttga 120
attcctctta attgggttcct gcaaaaattgt gttttttgg gttcggttgt gtgttgtgat 180
aatgc当地aaag gttgttaaaa atgattttta tttcattttt ttatgtgggg ggtttagtt 240
ctagggtgatt tgtatgaatg cacttggtct tggagaaaaaa aaattgtgtg aatttttagtt 300
ggaatggtcg aacagtgtga aaaagttta ggttgc当地t acacaactgt ggggtgtgatt 360
atgc当地aaag gtatattaaa caaattgaat tttatgt 397

<210> 773

<211> 468

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 773

ttgtcgtag aaaaagaata cactccaatt catcacatgg taaatgaata acaatggagg 60
acaaaanagta tatagcaata ttaataaaata aataaaaagt atatagccat tgaacgttag 120
tgacaaattc aatccatgtc agtagtcgca tgacattcct ttttcttcta gaactgaacc 180
aacatcacccg aaaaatcaaag ccattagccg aaagc当地ttt aacatagaaa cctataaaaat 240
tgtgagttag ggtttagtga taggtcatgg ttagttgggg tcttggatcc ggttcaagta 300
tggaccttcc tttccatttt ttttatcttgc actctcatga attaatgggc cagcaacact 360
tgatttgc当地 ttgatacatt atactaanaa ttagaaacct ataaaacctt ctctttctt 420
tctcttttcc ttttattatc tcttctaaat ttttcttctt cctctctc 468

<210> 774

<211> 444

<212> DNA

<213> Glycine max

<400> 774

aaagatctca gttttcttatt atataatgtat ataaatttgc atcagatatc 60
agttttctat tttataatga tataaactaa aatacctgtt tttaacactg tcttcatgg 120
gtaaaggcac agctccatca tcaggttctt cgtcatctt taccattacc gcttcatcta 180

ctattcctaa gttgtgaagc cattcttta acttcacaag acagcggtat tacatttagca 240
attaggctga atcagctata tcaaatgtaa tgaaggaaac taaactaatg gagatatgt 300
tcttcactta ataacaacgg agtaacatga taaagaatca gggtttgaa cctaagttga 360
agtgtgaggt tggggaggcc aaaaagatta aatgatagac acaatactgc aaaattaatg 420
tgcaaacctg ctcttgcttg tcat 444

<210> 775
<211> 456
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 775

aacccatgga agtcctaat atctcccaca cttgntggg tggccattc ttggatggcc 60
ttgattttct caaggtccac ttggacccca tttctaccaa ctacaaaacc taagaagact 120
atattatcta cacaaaaggt acacttctct atatttgcattt agagggtgtt tttcctaagg 180
actgaaagaa ctgcctgag atgttctaag tgatcatcta ggctcctact gtacactaaa 240
atatcatcaa aataaacaac tacaaatcta cctatgaaat ccattaagac atgatgcata 300
agcctcataa aggtgcttgg tgtgttagtg aagccaaaaa gcatcactat ccattcatac 360
acaccatact tggcttgaa agcgcgttcc actcatcact cttttcatac ctggattcgt 420
gataaccact tttaagatca tattttgaag agatat 456

<210> 776
<211> 442
<212> DNA
<213> Glycine max

<400> 776

gagctgaaca cacatacctc tataatagct aagcacacct cttgagaag agaagctaga 60
gcttatctac acaccccta taatagctaa gtcacccccc atgacaaaaaa acatgaaaat 120
aacagagaaaa agtccttatt acaaagacaa ctaaacatgc cccgaagtac aaggctaaaa 180
ccctatacta ctagaatggc caaaatacaa ggcctagacg aaggaataac ctattctaat 240
atttacaaag ataagcgggc tcatacttag cccatggct cgaaatctac cctaaggctc 300

atgagaaccc taaggccttt ccttggatct ctagccaat ctacttggag tcttctagcc 360
aatgcccttg cggggttaaga gtgcattcatt acttttcaact cagatgtgcg attcaggcac 420
atcagatatac gagacgctcg aa 442

<210> 777
<211> 286
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 777

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atgaactaag ggacgtcaat atggccaccg atgaagcctt ggaatgagaa accaagaatg 120
ccctgaagga agaacacgac caatacaagt tttgagggc tttatagggc aataatagtg 180
agctcatact ccgaagaggt gaaaggagtc atcacgggtc acaggtatga tctgtaaagga 240
cgagctatacg gcttgcccta tgacgagaag aaatttgcctt cgaccg 286

<210> 778
<211> 193
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 778

atcagaccac ttccagngtg ctggaactac ttcacactga tttgatggng cctattctag 60
ttgaaagcct tggagggaaag aggtatgcct atgtcgatga ggatgatatac ttcagaatta 120
cctgagtc当地 ctttatcaga gaaaaatcag acaccttga agcttaatt gagttgattc 180
ttatacttta aag 193

<210> 779
<211> 281
<212> DNA
<213> Glycine max

<400> 779

tcacacttac aaaggatata tgggtccatg agggacctcg ggcttctac agagggcttg 60
ttccatctct tcttggatcg attccttgc cagggattga tctcactgca tatgacaccc 120

tgaaagatct atccaagaga tatattctt atgacagtgg tatggtatta ctgcaaccac 180
attatctttt gaacttaatg gatttattt accactctga aatttttagt gacacataac 240
acatgtaaac tcaacccccc aacttaaata tgtaattttt t 281

<210> 780
<211> 247
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 780

ttcgctacta aagttggcga ctntgaaaat atttggagta gtttttaaca gattttatat 60
ggatatgtt gaatttagtgt actatattca ttcttatatat atgggcgtgt ctcaaatgaa 120
gtgcataat aagtgttcca gaaggttac gagtaaagaa aagcccacgt gattcccccta 180
actatccgc gttcatctcc acgtgactcc cccaaggctg caactctact actaccatgt 240
gctagaa 247

<210> 781
<211> 118
<212> DNA
<213> Glycine max

<400> 781
agttgttgg attatggcgc acccgtcata tgggtacta ggtggcgatc gggtgatgg 60
gcataatcaat tcttcacat ccacaataa gacatgaacc caccatcccc agttgtcc 118

<210> 782
<211> 260
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 782

tcagaccaaa gcaacatana atctatgtat ccgaaacccc tcaatcta gtgttgc 60
gggttgagaa gtgaatatga caatgagcgt tatttggagc aaactctcac ctcacacaag 120
tctataacat caatctaaac ttgctcaaac tggatttaca cctaaaactc cacccgaata 180
aaatttgatt cctcatcacc aattttaccc tagaaatgac tttcgatca ctacgtacat 240

ccttttcttt tattgcaaaag

260

<210> 783
<211> 449
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 783

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tctgatcatc atgctttgat aaatgcaaaa aaactggtgc aaatgaagag ggtgagaatg 120
agggagaaaac ccatgctgtg actgccattc ctatacagtc atgtttccca ccaacccaac 180
aatgtcatta ctcagccaat aacaaacctt ctccttaccc accaccagg tatccacaaa 240
ggccatccct aaatcaacca caaagcctgg ttaccgcact tccaatgaca aacaccacct 300
ttagcacaaa ccaaaacacc aaccaagaga tgaatttgc agcanaaaag catgtataat 360
tcaccccaat tccgggtgtcc tatgctgact tgctccata tctacttaat aagtcaatgg 420
tagccataac ctcaaccaag gttcatcaa 449

<210> 784
<211> 399
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 784

tattatgatt acgagctata aaaccaaact catatcta at gagtttagag ttaagttaat 60
tattttatat tttctctttt ttctgctcan agtaagat at aactttcaat tcatacttac 120
ttggctagca aaggaagaaa tcaattcttg cacagtcgat tttgtttat agatttacct 180
acatggagaa acaaaagtca atttgaardt tcattggatc aaatatttga taattggatg 240
cttaaatatt tctagcatta aatattcata aattgcatat ggcttctttt gactcctcaa 300
aaacataggt aacaaaatac attgcanatc caatgttagat tactggctca ttattatact 360
aatattgtta ttccaaacgta gattgctana tataatgt 399

<210> 785
<211> 333
<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 785

agctnggaag gatgcttcga tggaggataa gaaataggga gagggagaga gaggggggag 60
cacgaaattg aatgaataaa agagggagag aagtggact ttgaagtatg tatcacaaga 120
ctctcattca tcagagttac aacaagtgtt acacatgctt ctatgtatag acttggtac 180
ttcccttgaga agctttcttg agaaaacttt cttgagaagc ttgttgata agacttcctt 240
gagaagctag agcttagctg cacacactcc tctaataaga aagctcacct ctttgagaag 300
cttccttgag atgatcgtaa ataagctaga gct 333

<210> 786

<211> 200

<212> DNA

<213> Glycine max

<400> 786

gcttcacaat gacacaaggg agtgagactg tgactgagga gaggaggggt caggtctata 60
gggtgcgaga ctgaggagag ggaatgaaaa tctgtatgac gtgagagtga taacaacaca 120
tcatggccct ttacatgatg gtttaataaa aaccgatgtt gagtatctca ttgtcacaat 180
ggttatgaca aaaaacatct 200

<210> 787

<211> 365

<212> DNA

<213> Glycine max

<400> 787

tagtgcacacg agtagccatc ccattcttagc gagtgccat cttatcaaga tgctcaccca 60
caaatgctcg aagatcgctg agcttgtga ggactctatt ccacagagaa caacaagcat 120
gccttcgtgg tggaggagat ggggtacgta catcaagaac aggtggtggc aagtcttgg 180
agcgaactca ctgaccatta acatcccttc gataaccaaa ggaggttaaca acaccggcac 240
caatggagaa agaccttta accttgacat atggttcatc ctccaaagga acattggatt 300
gatgacgaca tagagtatca aggcggcgat acatgagagg tgcattgacc cgttatgcc 360
tatgc 365

<210> 788
<211> 117
<212> DNA
<213> Glycine max

<400> 788

agcttgcgg ccacgattga cgaaggcac atgatgacga cgtagtctc tgcgtttat 60
caagcttttc gtcttacaga tagcctatacg tttatacgga ctaccactcg ggtattt 117

<210> 789
<211> 115
<212> DNA
<213> Glycine max

<400> 789

ctataaatct aagcttaaca tcagaccctt ccaggtgtct gaactacttc acatttattt 60
gatgggcct atgcaggttt aaagccttgg atgatagagg tatgcctatg ttgtt 115

<210> 790
<211> 61
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 790

agctntggta cctattgatg gatacctaattt ttgttgatct aattnttagc attaacaac 60
t 61

<210> 791
<211> 405
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 791

agctncacgg cacgactgtt aggattaatg caatgattgc tgaactacaa gctctacagt 60
cgaatgagac ctggaggctc actcttcttc ctccacagaa aaccgccatt ggctgcaggt 120
ggatttacaa gatcaagtat cgcgctgatg gctcgattga aagatataaa gcacgtntag 180
tggcataggg ctacacgcag atggagggtc ttgattatct tgatacgttc tctcctgttag 240

catagttgac taccggcgt cttcttcttg cccttgctgc cgtgaatcaa tggcatctgc 300
ggcaactgga cgtaataat gcttcctcc acagacaact ttagtgaagaa gtttatatgc 360
aggttccacc gggattgacc gtttcacatc ctcaactggc atgtc 405

<210> 792
<211> 411
<212> DNA
<213> Glycine max

<400> 792

tcaagaaaatt gtaaaaagaa taaggtata tacatataca ttttagtacaa atagatata 60
ctttacacac tactgtatga ttgtgataat taaaaaagaa aaagaaaaac agttttcttc 120
tagtaggaaa ctaaattcat accttaggtaa agtttaaga atatctaatt acccatgcaa 180
aatttcagca accgaggcaa gatataaaca ttgtatTTT aggataatta gccactgaat 240
gtgtccttat tcttcgcaca gcaatgagga actaatccaa atgatatcg aaaatggat 300
aacaccagtg acaaataagac aatagtacta acagcaatca actacggaaa gttaaggaat 360
attgtatTTT aaggataatt agacacatgt gtccttattc ttcacagaac a 411

<210> 793
<211> 502
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 793

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taactattga aataaaaata agtgctatat taatgacctt tggggtaat acaaaataac 120
attactcaca catttaata ttttgtggtg gagaataaag gtggtcgcga ctatagcaac 180
ttctagggcc ggcgctacta ctggtatagt ataaaacttc cgaccattt ggccaaaggc 240
tcttcgctat gcgaaggat gggggagggta tgTTTatacgc acccttaccc ctgcataatg 300
caaagaggct ggTTTcgat tccaacccat gaccaacaag tcaccaaggc acaactttac 360
gggtgcacca gggctcgccc tcctactact tgcataatgat acttaacaga aattgcgc 420
tcagctgcca gcagaattca catagagaag tattattaaa ttagatggca tcaatataca 480

cagtctgatg agtcagcttg gn

502

<210> 794
<211> 454
<212> DNA
<213> Glycine max

<223> unsure at all n locations

<400> 794

ttaaatcttc aaaggtagtc cgtctgaatt caattcaa at cccttgccta ggattcacaa 60
gcaggctgct aactcttgct gat tggcct tagcaaagca aatcttgc at gccgaaag 120
actntatcct tcctctataa ccactggta ttccaaacga gagttgaggg cagctgagtc 180
aaatggaatc cattcccccc taacttgac ctctttggt gacttgtctt catgatcata 240
aacatttgca taaaactcct ttacaagtgc cacatctt cttccctt ccaaattcgc 300
gattntctta tgctagttcc tcctgtttaa ctcccttga aattcctcan agtcattgaa 360
cgacaattgc acattcctt caggaatgat cttctgccg aagatattgt ttgtgtaatg 420
atcccattgct tcanggaaag aaaatttg tcta 454

<210> 795
<211> 306
<212> DNA
<213> Glycine max

<400> 795

agcttgcgc tagagctgac ccatcaactg ccctaactct tttagactgg tgatccctag 60
gctcttgacc ttgacttgat agaaccttt tctaagcgaa ggcatttgac ttgatcccat 120
gttttactaa agtgaacaaa aatcggtgcg aatcaaaaact ccaacatcta tcatgggtgg 180
aatggatgaa tgcataaga aatgcataatg acacatatgc aatttatgaa tacgggagcc 240
cgggaaatgg ttccttctt agatacaacg tcttgggta acaaagcgcc caacgtatgt 300
atttaa 306

<210> 796
<211> 399
<212> DNA
<213> Glycine max

<400> 796

tacccatcac atgtggtaact aggtggcggt cgggcgatgg tgcacaacaa gttttccac 60
atccacaatg cgcgcataaa cccaccatcc cctgttgccc acctccaact gagctcacgt 120
actccccacgt agcccatatc ctcgttctc tcaacaccgg gtcccatca atcctccaa 180
gcttccacaa tatccaagca aaacaacatt cacacagcac aagctatcac agccaagcaa 240
aacagagcaa aggcagaaaa ctctgccaaa acaccaacca aaaatcacag ctttcccac 300
tcaaagaccc cagtaacaat tccttcgatt caattcgtaa accgttggat cgactccaaa 360
atattactgg aagtctatacg tgcataagcc tacattctg 399

<210> 797
<211> 417
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 797

gcttctgagc atcatcaaaa taggcatact tattgtgctc aggtaataat tntagttcca 60
aggaagggtgg ctgaacagtg gaaggcaagg gaagggtgac agttggtgga agagaaatag 120
aaggatctga acctgcacct gcatcacaat cagaattgca gaagaaatct ctacacatata 180
attacataca gtagtagact cataagaatc atcacaagta cgatcaanag aactaagatc 240
aaagtctaaa aaatcagaaa gtaaatctgt acaaatatcc acactatcta ttgcatcatc 300
aatgatatct ataaggaaaa cataatgctc atgtgttagga tgtctcatgt gcctaaaaat 360
gttgaardtgc acaacatcat caccaaactc catggaaaga gttcctacat gcacatc 417

<210> 798
<211> 172
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 798

gcagcttgct acgattattg tcagacnata attatttatt taaaataaaat gttgttaacc 60
gattgtcata gagtatgcta attagttata gtattatata gtcagggatg cacaatctc 120
tccccaaagca attttcatt tgtatctgct ctcacagcac cagtagtgtt gc 172

<210> 799
<211> 106
<212> DNA
<213> Glycine max

<400> 799

ttggcagaga gccagaaaca ataaatgatg acgttaagc taatattaga aagaaaaatt 60

gcaggaagcg aagtgatcct ttttatggct acataccaaa aacccc 106

<210> 800
<211> 372
<212> DNA
<213> Glycine max

<223> unsure at all n locations

<400> 800

gcacctgccc ctgcagctt gacagtaaat ggaatggtag ccctccatac atggtaatat 60

tattcaacgg acccatacta tgattaaaag acactgaaac tggntctcga aatgtgatcc 120

catttgatcg ttgcacatcgag attctgaatg ttctttccaa ttcttatttct tttctcactt 180

catcttccag gtctcgttgt tgtgcaaatg tcatactcgag agcaatgatt tctctcctga 240

ttccttcctt ctctagctct cgccatattt tctctttctc caactctcgat tggAACACTT 300

cattgacatt gatcggcatg gccattagaa aaactctagg cactgaaaca tttctggaa 360

agccacactga ta 372

<210> 801
<211> 438
<212> DNA
<213> Glycine max

<223> unsure at all n locations

<400> 801

nnggaaataa tatntaacaa ctgttagtatt gaacttaacc cacaacaagt ttcttagttat 60

atgaactaat atatgtgtca cttctaatgt ttctatggct gttactttta ggtaaagact 120

gggaagacca gcttggagca aatatcaaga gtttatttggaa atcaggtaag ctaaaagcta 180

atagtcttgtt ccattctttt ttcttatcca tgcaccttta tgtacttgag aatccctaaa 240

catacatgtt acaataattt ttccccatattt gtaaaataac ttgacaccct cgaacttctc 300

aaagtcatcc caatttctat tcgattcgcc attggtaactg gtattttctt cagattatgc 360

tcctgtcagt gtttgtaca atcaattgtc tggagttcct ctattattca tattccaat 420
atgaatattt aaaatgaa 438

<210> 802
<211> 451
<212> DNA
<213> Glycine max

<400> 802
atcctcttag tcacctgccg catgcaagct taaaattgac aacggaagct ctccagaatc 60
tcatatggtg ataacttac acacgaaagt ctgattcagg cgcatagttt atctagacc 120
tcgaaattaa acaacgaaag ctatcgagaa actcatatgg tcataaattt tcacacggaa 180
gtccgattca tgccataat atatcgagaa gggtggattt gaaccaccaa tgctctcgag 240
aaattcagat ggtcataact tttcaaacag aagtccgata tatgcgcata atatatcgag 300
aacgttggaa ttgaaccacg aatgctctcg agaaattcaa attgtcataa ctcgtcacac 360
gaaagtccga ttcaagcgca tactatatct acacgctctg aacttgacaa cgaaagctct 420
ccagaaattc atatggccat aacttgcac a 451

<210> 803
<211> 377
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 803

ggccgccacg gagttntccg actatgctct tgtgtgggtt aacaagctac aaaaggagag 60
agcaagaaat gaagagccaa tgggtgatac atggacagag atgaaaaaga tcatgaggaa 120
gcggtatgtt ccggcttagtt actcaaggaa cttgaaattt aagctccaaa aactaaccct 180
aggcaacaag ggggttgagg agtatttcaa gggaaatggat gtgctcatga ttcaagcaaa 240
tattgaagaa gatgaggagg taactatggc tcgatttctt aatggtttga ctaatgat 300
ccgtgatatt gttgagctgc aggagttgt tggaaatggat gatttgcttc ccaaagcaat 360
ccaagtggag caacaat 377

<210> 804

<211> 153
<212> DNA
<213> Glycine max

<400> 804

agcttgttct tgattattcc ttagttctgt aacttgctta gaacaataaa cttggccttc 60
tcttatttgt cttagggctt ggcgaccacg atcaacaaag tactttcgac acctactata 120
tgttgactcg accaacggcg ttattggaaa gtt 153

<210> 805
<211> 246
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 805

gcttcctttt aagtgcgtca ncgttaana accgagtctc tggatgggtt ggcgaaggct 60
ttgatggta cctcggcgaa aagtgaatgg nggaacatcg acacttcctc attcagaata 120
cngggccccca cactttgcaa tgggtgttat ttcaaggtaa tgggatataa tatccctgcc 180
tatatgcttgc cctcttggaa agaaccttgg attcatgccc ctgggagtgg tcccttcaac 240
gattca 246

<210> 806
<211> 368
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 806

actaatgttag ttnctaaaca aaaatcaatt gaggaagctt cgccaagttt ccccaattgaa 60
aaacctttat tcaaaccctt caaagttgtt gaaaaggcta aacgaaaaat tagggaaattt 120
agaaaaacta aatccttaat tgaaggcgta ggtgacaatc atagtgaatt actaaacaag 180
aatggtagtt tacttaaggt cattccagat actccccaaag cctcgaaaaa tacttccaaa 240
atggtaacaa gaagtaccc caaattaattt aatatttata atgaagatag tgacccaaac 300
tcagataaca caactgagat aggatcagtgc tcagaaaaga atataaatcc aattaatttc 360
aaacactg 368

<210> 807
<211> 223
<212> DNA
<213> Glycine max

<400> 807

agcttgagat gaggaagtgt tgaagggtga aactttctgc ttttatttgtt gaccacagag 60
tggtaacctgg agatatgtcg tgggggtcac gagaccttgc ggacgtcagg tggtgtgcta 120
ttgcccaaaaa ccaagcttga ccaatcccga cccaaccgg gcatagtcgg taagtgagaa 180
cctgtatgtt acctaaacag gcgagtcctt ggcagtcaac aga 223

<210> 808
<211> 427
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 808

gaggtccagg aagggttggc ggccgaagga actagttccg ctccggagta cgacagtcac 60
cgctttagga ggcgtgtaca ccatcagcgc ttgcaggcca tcaaggatg gtcgttctc 120
cgggagcgcac gcgtccagct cagggacgcac gagtatactg attccagga ggaaataggg 180
cgccgacggt gggcatcact ggttactccc atggccaagt ttgatccaga aatagtcctt 240
gagtttatg ccaatgctcg gccaaacagag gagggcgtgc gtgacatgag atcctgngta 300
agggtgtcagt ggatcccggt tgatgccac gctatcggcc aactcctagg atatccngtg 360
gtgttggaaag agggccagga atgtatggcg cctactangc accctttgga cccagataag 420
tncaaca 427

<210> 809
<211> 361
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 809

agctttgata tggaaattaa gttganagta aatgtatggta tgtataagac atcaatcagt 60
gtaatgaatg cagaaaagttg tactgcgcct gagtggttag catagacaag atggccattt 120

ggtaatctaa ccgtgatggg attaatttga tgatatgagt gaaagttgt taaggaggag 180
gaaacgtgat cagtggctcc tgaatcta atccaggagg tagagttgc ttttcgtaa 240
gataggatta tacctgttgc atcgttattt gaacaagata aaatggaagc gacctgtgg 300
ttggtgatg ctgagttcc agcagatggc tggatgttata atgctagcca tgccttgtac 360
t 361

<210> 810
<211> 360
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 810

ntgcggatat ggtctttca atgaaaggat caaagtgggt ctggaaaaag gcaaataatga 60
tcatccttct tggatgaatg agaaaactgg ggcaaatgaa gaggatgaga atgatgaagg 120
aacccatgtt gaggctggca ttcctacaag gacaaacttt ccctccagtt caaaggccac 180
ctatattaagc ctgaaatcag aaatagaagt ggacgttggg ctttccttg agctttgca 240
tttttagata ttctataga gagaaaggc caagttccaa agagtttga gagctttgt 300
tgtgcgaaga ctgacagaga actgagatg aataggaact cattctgaga catgagatga 360

<210> 811
<211> 462
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 811

tgtatgtac aacttctgan atcaagaaac acacacacac acacacacac acacacacac 60
acacacacac acacacactt ttccctagtc gatcactcac ataaanttcc attctcccc 120
ttttgtttt aatttatgct tatcttanaa ttaagttat tactcatgtg agttcttgat 180
ttaatcccta ttctctcac ccttggcat caacaaaaag ccaaagtgc tatcanattt 240
gaagtattca aatataacta aatatccata caacattcat ggaaaaaaaa tatcaaccaa 300
atcatgaagc aagaaccatg aagcaacaat tatgaataga ttataaaatc cacatagtca 360
aacaacatac ttaatatnng ttcanatacc ataataatat agccaaaata caaggctgaa 420

gatcagagta ctaataatat taanatagac atctaagatg ag

462

<210> 812
<211> 297
<212> DNA
<213> Glycine max

<400> 812

tggcactgca gatctgatct gcgtgatcct catccacgta cattgtgtat gctccggttg 60
agtcgggtgta ccccttcttc gtgtacacaa cctcggtgct aaccctgctc ttgcattgca 120
acataatctc agcacctgca aacaatcatt aatcaaacaa tacgtgttac aaactagcaa 180
caactactaa ggtaatgttt ccattcacat taaatcattc aaaattacct tctaataatgt 240
aaatttaaat atatgttcac atgttagtc tcattgcacaaa aaaatgatgt tagatct 297

<210> 813
<211> 420
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 813

gcttcataata tatataacat aacattatgt ggtaagaccc aacacgcca caaattattc 60
tcaatttagtt gatgaacttc caagatccca tttcactaaa tttccttagn tattagatac 120
ctctaaatat atgacaaaggc atggatttga atatataat aagctcatcc ttatctttgg 180
gatccaccc agaatcaata gaatattcaa aatagtgtt aaggagcaacg aaaagaaaact 240
aagaacaata taaaaaggaa taaagtgcata accatctcac tcaatgaatg aatgctntga 300
caacctcaat ggatgcagac aaaataatca gaaattctag tactaatgtt tcttttagcag 360
cttatggta cttgaatgga gaataagtgt ntttctgag ctatggctc aacttgatat 420

<210> 814
<211> 432
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 814

tttcattntg ctattnccct tannattttt aattntcatg ttgtatatt gtcacgaatg 60

gacagtttt agtacattt ctttcttnt tctgcttgtt attattaaat agaattttt 120
gttgagagga agcagagagg ggtgttatcg ttgagaggat gcagagaggg gtgtggcaaa 180
atgacgttaag cgggttcctt ttacaatcta tttatcccac agggtctatt tttaaacaat 240
tgtcgaagag ggtctgtttt tttaaagggtgt tccaccactt ggactggcgg aacccttgct 300
tgcataaatt cccatgcatg catggttctg ccagtgggtgt tagcgtaaat caacccaaag 360
ggttcgcca atcccactag tgagataccc catgccattc atcccatcat ccatcaacgc 420
catccatcca tg 432

<210> 815
<211> 429
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 815

ctaaaccccttg aaaacttgtg ctattcattc ttttcatctc ttctccctt gccaaanaga 60
attcgccaag gacctaaccg cctgaattct tttgtgtctc tcttctccct ttccaaaag 120
aacaaaggac taaccgcctg aattcttttag tgtctccctt ctcccttgta aaagaattca 180
aaacgacaca gtctgaaaat tcttttgatt ctccccattc cctaatacaa aagtgttcaa 240
aggactaacc gcctgagaat tctttgtat ccccattcac aatgtatcan aggttaaca 300
gcctgagatc tttgtctaaa cacattggag ggtacatctt ttgtggtaca agtagatgg 360
acatctactt gtgttgact gagaacaaga gaangtacat ctcttgtaa tctgttctag 420
tggagggtta 429

<210> 816
<211> 529
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 816

cgtttttttt gattgaaacg ccatctatnt angtgtgact acagataact caagctgtac 60
ggtattgcgg gcaagttatg tgctgaagat ntattnata aaattaatta attaattgat 120
gtgagttaaa ttgaaataa aataaagaaa caggacatcc atttaccaga acgtgtccca 180

attaaatttt gttactattt ttaattacat aaatgatata tatatatata tataatata 240
tatatatata tatatatata tatatatata taattggcat gaatcgaaaa tattctttt 300
ttcttttta tcgtaattt aaactgtaaa aaagttaggac attttttcc tatagcgtga 360
tacatacaaa gttaacagtt aacaattgtg tntatataat atataaatat atattaatta 420
ataatatata ttacttctag taaaaataaa ataataact gcttaataga tntgatgatt 480
aattccatcg attatatact aatctntnta atataaaaaa ataaaaactn 529

<210> 817
<211> 501
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 817

agtcacctgc ggcatgcaag cttctggttg ctgcgatgat cgaccagaac tgttccctga 60
actcatccaa gtcctcaaag gcttgagact cacagcagtt aaagctgaca tagccagtgt 120
tggtggcaga atcaaaagca tattggtgct ttgttctaag gatagagaag acagtgttg 180
ccttgccact ctcaaacagt ccctcaaatc tgctgtcacc aaaattgctt catcatccat 240
ggcttctagn tgcggcgta gaagtaagag gcagagattc ttcttgcctt ctcactgcct 300
acagtttaatt atttattgca aanaatattt tttccccac tattcattgc agtatgggc 360
aattatttgc tctattnca atatatatat atatatatat agactccccca ttaggaagat 420
aaaatcatga aatatttagtt ctgtgcaacc aattaaggca tagttaaatt ganaggaaag 480
gacgcacagc atagtatgag t 501

<210> 818
<211> 447
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 818

cgcttaatta tcttagagg aggataccgt ccaatggagg ataaaaaaat gatgttgata 60
atctttcct ttttacatt tgaaggtatt aaaaaaaaaagt gtcttcatga cattattaag 120
.aaattaaaaa aattattaaa aatcatataa tgcatttaaa aatcattgga gaagacattt 180

ttatgcattt caatgaaaat atttttcttt ttatTTTTT taaccgctgt tcttaaaaca 240
ttagttataa ttaccttaa aaaatgctaa caaaagatat taatgccttg ttaatttagaa 300
gtattaaaat aaaagattga ttctttntaa aatgtatata ttgtttataa tattttattg 360
ataatacttg taatttagtgt cacaattaaa agtagcttt tataaaaatt gaaaatgaaa 420
acttccattt gaagtcaGca atgttat 447

<210> 819
<211> 429
<212> DNA
<213> Glycine max

<400> 819

tTCGTTGGCG aaaggatcta tgtgggtctg agaagaggca aatttgatta tcctgcttta 60
atgaatagga agcctgcggc aaatggagag aatgagaagg agggaggaac ccatgttGtg 120
actgctgtcc caacacgacc aaatttccta ctGctcaac aatatcaata cttatccaat 180
atcagccctt ctcattaccc accaccctat caaccaagaa cactcaatca ttcacaaagg 240
tcatacctaa atcagccaca aagcccgcct accgcatatc caataccaaa caccaccctt 300
aacacaaacc acaataccaa ccggtaatg gaatttctag aaaagaagcc tgtagaattc 360
accccaattc tgggtttgtta tgctaactta actccatatc tactcaataa tgcaatggta 420
gctataatc 429

<210> 820.
<211> 466
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 820

ntccctcat tctcacgttg ctTTTCTTCT cccCTTCTca tccaccattg ttGCCCTTta 60
aagctccaaa ctttgctcac catttctact ccaaattcaca aaaggaagcc atTTTcgag 120
tcgtgaagcg cacctctacg ttgtggact tcaaattca ggTTTggta gacttcttct 180
cacataaatt ttgtggat tgggtctttg ggagatatga tggtagttc tactagttt 240
atgccttatg gtatTTTTT gtgaaggaat ttgtgaaag catgctaaac tcgtcatgtt 300
tgatgtgagc caaatatacc cattctgttt tagggtttta taatgtatgtt 360

tgtgtgctga aatcatttgt agaaaactgg tagagatgat ggggagagtt aacctanggt 420
taaaagttag aatggtagtg atgtgagtgg aaaagtgagg ttttga 466

<210> 821
<211> 483
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 821

gcttcacact tgataatgga gacacatgaa cagtgcatac taatgacatt catggcgctc 60
caaacaaagg tggagtatgg aggattgcct tgagggtccg cacttaggca atcatgaaac 120
tcaactccaa actcgaaagt ggaggacaca tgaacagccc taagcaataa cattcatgtg 180
gctccggaat agatgagaa tggaggattg ccttgagggt cctctcttaa gcaatcatgg 240
aacacaactc caaactcgaa agtggaggac acatgaacag ccctaagcaa taacattcat 300
gtggctccga agcangatga gaatggagga ttgcctcgag gtcctctct tatgaaatca 360
tgaaactcaa ctccacactc gaaagtggag aacacatgaa cagccctaag caataacatt 420
catgtggctc tggAACAGGA tgagaatgga ggaatgcctn gagggccctt cttaagctat 480
cat 483

<210> 822
<211> 412
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 822

agtctcacga ttgtcacgtg ctcacatgaaac atttggtagc cgtggctata tgagacatct 60
tgccaaacaa agtcaggtta acgataactc gcctgtgctt tttcttccat gctatatgt 120
gcaaagtcat tgatccagtc atgtttgatg atttggaaaa tgaggccgca attatactgt 180
gccagttgga gatgtatTTT cccccctgctt tctttgacat catgactcac ttgattgtgc 240
atctggtcag agaaatcaaA tggtgtggtc ctgtttatct acgggtggatg tacccggttg 300
agcgatacat gaagatctt aaagggtata caaagaatct atatcggtca gaaacatcta 360
nttggagag gtacattgca gaagaagcca ttgaattttt ttcagaatac tt 412

<210> 823
 <211> 453
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 823

agctnttagga gaaaccatat aaactaaggt agttcctana caaaaaacaa ttgaggaaac 60
 ttcgccaaga atccccattt aaaaaccttt attcaaacctt ttcaagttt gtgagaaggc 120
 taaaagaaaa attagggAAC tttagaaaaAC taaatcctta attgaaggcg tagtgacaa 180
 ccatagtgaa ttactaaaca agattggtag ttacttAA gtcattccag ataccccccA 240
 agcctcgaa aatacttcca aaatggtaac aagaagtacc tccaaattaa tcaatgttat 300
 taatgaagat agtggccaaa actcagataa cacaactgag ataggatcag tgtcagagaa 360
 gaatataaat ccaattaatt ccaaacactg gagaacaacc tccatattat attatcaacg 420
 tccaaactggc cctgaccttc tattagagga aag 453

<210> 824
 <211> 457
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 824

ngaaacaact cgttacataa tttgatatgg ctggcaac agatttcata agtatttcgt 60
 tacccattct agaaagagat tgctccatcc acttgttcaa ttttattac acactttct 120
 aaatgctaga gaaaatcatc taaaacttcc ccaataaagt ggtagccaa gatatttgc 180
 atgaaaatttgc taacgacctg tcttgtcgat atgatattcac cactctaaag tacgtaaattt 240
 ntaattttta aatgaaaattt tcattaattt gcttatgaaa aatgagagta aattttcgc 300
 gatatagatt caccaaacaa cgcacaatta tttaaatgaa atatataat atatataat 360
 atatataat atatataat atatataat atatataat atatatnaac tttagccacac 420
 tcacataata gaaaagtaaa tttagttcata catatag 457

<210> 825
 <211> 479

<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 825

agctngaaca cctattagta tttatatttc ttaaataata aatgtatgac atcaactcgt 60
cattggctcc caggttagt gattaagcaa aatagaccaa tacaaactca cgggttaagt 120
cacctaacc attgatccaa agtttacatt gtcacctcta cattagtgac tnttggtgc 180
ctttgttcc tttaagcttt ntgtgtataa aaatatattt tttcttgtgt gaaatatttg 240
tttggaaattc agtttaact atataataaa attgatggtt aagtttaata tatatttaaa 300
cagtcggat catggat tgaggacttg gataaaatat atattttca aagttttgtt 360
aatataactt ggtaaatata attctaattt tataaactat gaaaaatac aaaagttaga 420
tgaattcaag ctcaacacaa tagaacaagt accaakanat actatcatac atttgacat 479

<210> 826
<211> 430
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 826

tctgtctgca gtggcaccac caccagccat gtttaattat ttgctctcaa acaaccaatt 60
aattctctag cctcaacgatc ggtacgagct taatttagtat attaagaatg ctattattag 120
taatataatat tcagttctat agagaatgat gtttgcatt atgcttacag accgtaatgg 180
tattgctctt gcgggaacca catacactct agctagaaaa caagacatac atagttaaatt 240
aattaataat gttaaagccg gcccttgaa gacaatttac atgcttaagt ttcacgggtt 300
taggtctaaa taatgccatt aaattattn tttgtttgg aatataattt atttattatc 360
atttaaatgg caacaaggct tggtcgttac ttgtttgttc tattatgcgt ggagtcttat 420
tcatcgaaacg 430

<210> 827
<211> 309
<212> DNA
<213> Glycine max

<223> unsure at all n locations

<400> 827

agcttntgac cggctgacaa gcaacaatct aagttataat atctacacca caagctgtat 60
cattgtaata acttactcat attcttcaca tttggacaga cgataatatc tacaccacca 120
gctgtcctcg aataacctga attgaaacag aataaacagc agtaatggaa taaaccccat 180
cgaaggcatct gcgtaagaaa cccacttgg a ggtgtatca aatatcaaaa aggaggctgc 240
tgtaaaagcc acaaccatat catcaacata gataaaaagt aacacggtcc caaaagtcaa 300
gctcgatg 309

<210> 828

<211> 222

<212> DNA

<213> Glycine max

<400> 828

cttctgaacc ttccctaagga aatccttgat aggattgaga ctcaggaaga tatgaactat 60
attgaagggtt ggttagcaaat cttaacttct acattaacgc ttctcattat gtataggata 120
cgagtgaaaa ttcatatctg ggatgggta ttccggatga atctttcttgc cccgcgttcg 180
tcattcgatg cccatataat accctcatta aagggccaca ac 222

<210> 829

<211> 406

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 829

cccgctgagt tctttgttat caccganact ctaggaggag gccaaagaac tactgtgtaa 60
gattttctt attcccttgn tagtgttctt gatttgtgaa tctcacttaa attttgagct 120
taatatgtgg catgcattgt gaatcacatt ttaatctt atcagctaag ttgagttgtt 180
tatgtatgtt gtagggcatt tcaaggagaa acgaagcaat gagcttatat tctaatacgct 240
canaatcaca tataattctc acatttgc ttagtctt gtgttaaggaa ctgtcaaatt 300
ttgttaattct acctaacatt accagcagtt gtgtatggaa attgtntgtt tcctaagatt 360
caagccaggt ttatatttc tctgttagtt ctattgctaa acatga 406

<210> 830
<211> 399
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 830

gggtgtacgt aaaataatta gcactaaagc ttatattatt ntatgtattt atcacaatgt 60
gcaaatgaga tgattgtgtc taacttttc cataacatat aggcctgatg ttttatttta 120
gaagttata atctgatctt cagaatttga aatttttct ttacattctt atcaaaggaa 180
cgtcgttta atcaggtaga gtacaattat aaacatataa attacaaaaa ttaaatataa 240
catntatgg tttatgtga tgagtgagat gtggtaatgg gagagcaata aggaaagcat 300
tntggatcca canaacagga catgaagctt cttgcaagtt ttcaaataat cccttcacga 360
atcggttgg gatcttgca aatttcttgt ctcanatcc 399

<210> 831
<211> 440
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 831

tgcctaagtc agcaacatac acactgtgtt tattgtggat tctacaataa aaggttaactt 60
tgaataacttc acagtcatga agtgtctacc aatctctgtt aagatagttg ttacaacttt 120
attnaagcaa agaaatagct ccttcctta aacacccaaa cagaagcaac aaaagtttagc 180
aaagcaagaa atacaaccgc gaaccataaa atagtgtct tatagattca tgattcttaa 240
tattttcagg tgtgttaaaa acgattttc tgaaaaattt gtcacaaaatc aagcttgaac 300
cagagaatca ccacgcgcag ataattaact gcacattagg tgccgtagct atcgagataa 360
agaaaaatgg taggttattgc tcgagaagaa nagagaaaa gcttatgaac aagaaagcan 420
ataaaattgag caaaatgatc 440

<210> 832
<211> 313
<212> DNA
<213> Glycine max

<400> 832

tctcctacca ctgccttaca atagtcatca agcaatatgt tggcagccctt cacatctcta 60
tggattatct ttggatcaca ctgctcatga acgtatatta gctcccttgc tgctcctaag 120
gcaatttgct ttcttgcgcc ccagtccaaac actggcttac ctgcaaataat ccatcaactgg 180
tatatggatc agccactcaa cttacatagc acatgcatgt tcattctagt tctggttaca 240
taaacttctc tatatgcacc tttgagaaga acataagaag tttagaatcaa tttatgtacc 300
ttagttattg gag 313

<210> 833
<211> 336
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 833

tgtacgcaca tcgttcgcgt gtatgatatt cactccacaa tgttcgaagt agaggagac 60
ttcaacccta taacgcaacg tggcagacaa aagtggcag taaacttgaa tggtcgcat 120
tgtcaatgcg gaaggatttc tgcgcttcac tatccatgtt cacacattat tgcagcttgt 180
ggttacgtga gcatgaacta ctaccaataat atagatggtg tttataaaaa cgagcacatc 240
ttaaaagctt actccgcaca atggtnct cttacgaatg aagcggctat tcctccttct 300
gatgacgcattt ggacacttat ccctaaccac actaca 336

<210> 834
<211> 252
<212> DNA
<213> Glycine max

<400> 834

agcttgtctc gctaagcgat aatccacttt tggctctaaa cacgactttt cgcaactaagc 60
acaatttcct ctcgggttgg gatttgcgtt gagtgtgaca attgatgttg agcacaattc 120
atttgcgtt gagtgcaata attcgcgttg agtgcaacct ttcatcccga gagcaattcc 180
ttcttgggtt ggaattgcgc ttagcgtgct tctcgtgcta agagagatgt aaaaaattgt 240
tgttctaaat cc 252

<210> 835

<211> 254
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 835

tgagatgagg aagtgtagaa gggtgaaact tcctgctttt attcggtgac cacagagtgg 60
tacctggaga tatgtcgccg nggtcaggag accttngga cgtcagggtgg ggtgctattg 120
ccccaaaacca agcttgcacca atcccaccc aaccgggca tagtcgtca gtgagaacct 180
gtgatgtacc taaacaggcg agtcctggc agtcaacaga taaaaggaac aaagaccaca 240
aagcatggag gctt 254

<210> 836
<211> 402
~~<212>~~ DNA
<213> Glycine max

<400> 836

tgcagcttgg ttaatattag gccttgataa tcaataaagc ctattggaaa ttaatatgt 60
aaaggttagt actaaaaatg caaaattaca cttagttt tcataagcat aacatccaat 120
acaactcata agtttataat tagtcacata agttttcat aacatatcac aagtcacaac 180
taaaataaaa gaaaacagtc caagtgtgca attatagaat ctagaattct tgatatttag 240
actagcacca aatcgctaattttccaaa taaaaacaca aaggtagtaa tagagatgg 300
gagggcact tcatgtgaga gaaaagctt tctctgctaa gcctcaggcc aaagtccaag 360
ggtgaatatg gagagatcaa aagttacaga gcaatggta ta 402

<210> 837
<211> 433
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 837

tgattnatgtt ntgcgtgacg aaaggatcga agtgggtctg ataagaggaa aatttaatta 60
tcctgcttgc tgcttggacg aatgagaaaa ctggggcaaa tgaagagggt gagaatgagg 120
aaggaaccca tgggtggct gccattccta catggacaaa cttcccttca gccaataat 180

gtcatcgctc agccaatatc gacccttctc attacccacc acccagtcat ccacaaagg 240
catccctaaa tcaaccacaa aacccaccta ccacacaacc aatgctaaac accacctta 300
gcacaaacca aaacaccaac caaggaaggg aatttgtagc aaaaagcctg tagaactcac 360
cccaattctg gtgtcctatg ctaacttgct ctttatcta cttgataatg caatggtagc 420
gatcaccct act 433

<210> 838
<211> 111
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 838

agaggtaaat gacatatgtat catttctcct acttgctgtt gcactttgg ctattttga 60
tacctaaagn tttcattncg aatttaatta cacgatgatg atgctggatg a 111

<210> 839
<211> 402
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 839

agctngcgaa taattaacac agaatngtac aaaattctta tgatacatga tcaaatttat 60
caaaaaanat aataatgacc cctgaagcta tcttggtgac agtgacaata agttgagcct 120
tgtgcagcaa aacttagtgt tgagtgaagg atgacttggtt gcttgtgaat tgacttaacc 180
agttttgac agcttaccc ttggcaatga agcagccatt gttcttccta tcaaccttca 240
acggactccc catttgacc atcatttct gaaccaacac tctcaccggc gactctcctc 300
tgacactgct atgctttccc aacccacana ccaccgtaaa ttccgcagga actccataat 360
tggaatcatt caacccctt ccatctcctc aaccaaagca ca 402

<210> 840
<211> 116
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 840

taatacagta gtctttatcc actttcaaatt tntttactg ttatgggttc tgatttgatt 60
 ntgttttggaa ttgtatccggc ggagtttgct ttgtcttaa ttattgatta tttgat 116

 <210> 841
 <211> 303
 <212> DNA
 <213> Glycine max

 <400> 841

 ccaacgccagg ctctgaccac tgttcttctt ttccgcgatg cttctttca tgtccgcccgg 60
 agtgggctta tagcctacac catacttgcc acgattaccc tttgttttga tcagacttagc 120
 tatggcgcca ttgtcttgc ctaaacccat tcgcgggtca taacgggtgc tcaacatcac 180
 tcgggctctc attacctcca ggtatgaccg acctgctgca gcgacacgtg cacctcagga 240
 ggaaatgctg accaccta aagactggag agcgggtctt aacgattctt ctgcgggttcc 300
 aca 303

 +

 <210> 842
 <211> 289
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 842

 agcttcaaca natgttcacc agaaatccta agctatcacc atcaacctct aatgacatta 60
 gtttctttt ctttgttntt cttccataat catttcctt ttcattctt cattgtgtgg 120
 tgaactcctt tccatggta agcatacact gcaacaatct ttagcccaag atatggagtt 180
 tgattctaaa cgtaagttga cttttctta ggaaaatatg cttcattntt tataacaat 240
 cttgttctgt tgcaacatac attacaacct ntatataata tatatatata 289

 <210> 843
 <211> 436
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 843

 tcttagatag caacttggtn ttttgcttagta atgcatttttgc agagggaaagc tccaggaagc 60

ttcttttgt gggaatatga gtccgatcac acaagatggc atgatcaatt gcagccatgt 120
tctcaataag ctccatggct tcttctgggg tcttcaattt aattttccc ccagcagaag 180
catcaaataa ctgcttggac tgtggccta acccatctat aaaaatgttg aactgaattg 240
gttctgaaaa tccgtgagtc agtgtttcc gcagcaagct atggaatcgt tcaagtgcct 300
cgcttagaga ctcatccaga aactggtgaa atgaaaagat ggntgcttt ccttctactg 360
tcttagactc ggngaaatat ttcttcaaac atttctccac caattcatcc catgtcttga 420
gactgtntcc cttaaa 436

<210> 844
<211> 434
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 844

agcttcatga ctagacatga cttctatgac aaaactacaa taggtggaca agtcgctcta 60
gatttgcag gtttcttct actttaatat ttttgaaga attttatgtat ttaggtttca 120
gccacaaaaa ataacaagac aaaactcana tcatttggtc atgagtgtat gaaattctt 180
tagcctatta tttgatttga gtcaaatctt tcatttgcatt tagccttaa catgttcatg 240
caaaatgctt agagagtctt tgatttgaa ccttgctt aactttatg cttcctttag 300
attgcgtcta ttgtgaatat gagtcttggt gattgaattt ctggctgaaa tggtgatcct 360
aagtgaatat tgaactccta taactgtcgt aaacagtctt agtgaggtaa acatacatat 420
gaagggtgaa agta 434

<210> 845
<211> 376
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 845

ngaagctcaa ggaaaagctt gaagaagttt tggctattac atgccaact ctcttaagt 60
gcatttgcattt tggttgcattt ctnggtgtt tcttcttagt acatttgata tttgtattgc 120
atcatgcattt atcatggttt gtgtgaagaa aagtttctaa gttagaaaaa tttcttcaga 180

ggcaaaaaca ctatTTTaaT cgattacaac cttattgtaa tcaattacga caagctgtct 240
gaagcttata gagttgagtc tcgttatcaaA ttaatcgatt acagctatct cacaattgat 300
tacattattg ttcgagacaa tgactgattt attcaagagt ctctgcTTTA atcgattact 360
tctttctcgt ttaagt 376

<210> 846
<211> 415
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 846

agagatacaa tctactcttg gatttgcatt aaaanaaaaat cttgcattct atctttcttc 60
tgctccctgc ctgaaaacat gcatgtatag cctggtaatt ggtcaatgtc tttgctaaag 120
attatctggg ctgttgactc gaagctttc taatttattc aataagatgc atgaataatt 180
taatcataaa tcataaaattc catacatgat gtaattattc atgtatctt ttgaatatat 240
agatctttag gttatttcat ataatttagtt aattaactgg tgattatTTT ctgaccaagg 300
ctggtgatta ttccatacgc ttacgtaatt aactgattct gtttatattt ttatttatta 360
attcttcata atggagatga antctacaca attcatgttt gaacacagga tgcatt 415

<210> 847
<211> 455
<212> DNA
<213> Glycine max

<400> 847

tctttgagaa aacttccttg agaagctaga gcttagctac actcacccct ctaataacta 60
agctcacctc cttgagaaggc cttcttgaga agattcctat agaagctaga gcttaggtac 120
acacacctct ctaatagcta agctcacctc ctgagatga aaagctagag cttagctaca 180
caacccctat aatagctaag ctcaccccca tgacaaaata catgaaaata caaaaaattc 240
cctactacaa agactactca aaatgtctcg aaatacaagg ctAAAACCCt atactactag 300
aatggccaaa atacaaggcc caaacgaagg aaaaacctat tctaataTTT acaaagataa 360
gcgggctcat atttagccca tgggctcaaA atctacccta aggctcatga gaaccctagg 420

gccttcctt ggatctctgg cccaatctac ttgga 455

<210> 848
<211> 349
<212> DNA
<213> Glycine max

<400> 848

gcgagctctg accactgttc ttcccttcgg cgatgcttct tttcatgtgc gccggagtgg 60
gcttatagcc taaaccatac ttcccacgat tcacctgggt ttttatcaga ctatgtatgc 120
cgccattgtc tttgcctaaa cccatcccg gttcataacc ggtccccaaac ataactcgaa 180
ccatcattac cgccgcattct gacagacaat gttgcggaaa gagggaaatcc acggaggaaa 240
tgctgaccac ctcaaaagac tggaaagcgg tttctaacga ttcttcgtcg gcttccacat 300
aaggcatgga ggatgggcag cttaccaaga tatcttcctc gcctgacac 349

<210> 849
<211> 106
<212> DNA
<213> Glycine max

<400> 849

gctgatgagt actattgtgg agatgattga ccattctcaa agccaaggaa catatggatg 60
cgtattttga aatgatgaga tcattatcca acaacatgaa tttatc 106

<210> 850
<211> 358
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 850

agctttgcca aataaacaag ttctctattt tgatactca gatcattcca ttttaatttt 60
aaataacttgg cgacccgatg cgcttgcgg tataactt ctgctttgat gtaagtcttt 120
gtaaaattaa gaaaaaggaa ctgtgtgggg agacgaacag taccacattt catttgagag 180
ttgaggtcag gtacatatata cataactaagg atgagtgatt gaaactatgg acgaatgtatg 240
actactctgt gaggatgtatgt tggactaatg gatggttgcg tatgtttatg ggatctgata 300
atgttttctt actaattattt cgagtttgtt attaacttct tttataataa actcaccc 358

<210> 851
 <211> 435
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 851

tataccttatg gcttgccttc ggacttcact ccccgatcca ccccgaaaga tttaagccaa 60
 gcccctactt tcgagggca gctcccaccc tatgacgact atcccggca agacgatgag 120
 gaaggagata cccatctcggt tccccgtctc cacctcaaag atctgtcccc ccatgaacta 180
 ccccaaccaa acatagtccg ccataatcccg acttcaccca cactcgtaaa agaatctgtt 240
 cccttcgtgg aacataaggg aaagatttag ggcgttgaag agaggttgag agcagtcgag 300
 ggcctcgaaa attaccattt ctggatcta gcggacttat gtctcgtaacc caatatcgta 360
 attcctccca agttcaagt accggacttt gataagtaca aaggatgac atgtccgana 420
 gggcatcttc ggatg 435

<210> 852
 <211> 187
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 852

agcttgaata ggtgtcggt tctactgaac tatatatatt ntggtggnnt tgcaatgttg 60
 ctatcctaga ttttgttattt tctctttgg atattttagc gtcgacatag tgtaatatac 120
 aggtatgagc ttatcaagat gaatcattt agggattat cacttggcat tgttcagcta 180
 ttctctta 187

<210> 853
 <211> 333
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 853

nttaggtgga gcatattana ataaaactaag ttcatagaaa ggtataaga cacttcata 60

agccccaaatg agatcttcta aacaccctct ttatccatat aactctcaa gtattactta 120
acaaaaatttc aatgaatcct tcaaataatgg gaactgtcat ttcaatcaa gaaaccaaca 180
aagtccacca taaaaaagct gtcttcactc tttagactgt tttacaaatt cactagaaaa 240
tattctacga agaagatatc atcatatgat gtaaagcgtg ctagttaact atgcaatgac 300
acaaggcgtc caaatgcacc cattttaaga tct 333

<210> 854
<211> 315
<212> DNA
<213> Glycine max

<400> 854

gatgatgcag atgggttgt agtacctca tgcactcctc taatgactat ggcattcattt 60
ctggcactaa actgctggga gttggaggcc atcttctcaa ttaaaatttct ggcttcagca 120
ggagtcatgt ctccaagggc tccaccactg gcagcatcta tcatacttct cttcatatta 180
ctgagtcctt cataaaaagta ttggagaaga cgctgttctg aaatctgatg gtggggcua 240
ctggcacata gttcttaaa tctctcccag tactcataca ggctctctcc actgagttgt 300
ctaatacctg agata 315

<210> 855
<211> 303
<212> DNA
<213> Glycine max

<400> 855

tcagaagaaa gtgatgaggt acaagctcta aaggcagagc ttgaaagagc ccgagtagtc 60
gaagagaagt tcaagtccat agccatcaa gtcgaaaag agtatgatga actaaggac 120
gtcaatatgg ccaccgctga agccttgaa cgagaaacca agaaggcccg aaaggaagaa 180
cacgtcaag caaagtttg aggggctta tatggcagca atagttagct caagctccta 240
agaggtgaaa ggaatcatca ctggtaaag gcatgatctt gaaagacgag ctaaaggctt 300
acc 303

<210> 856
<211> 415
<212> DNA

<213> Glycine max

<400> 856

cccatcacat gtggtaactag gtggcggtcg ggcgatggtg cacaacaagt tttccacatc 60
cacaatgcgc gcataaaaccc accatcccct gttgccacc tccatctgag ctcacgtatt 120
ccccacgttagc ccatatcctc gtttctctca acaccgggtc cccatcaatc ctcccaagct 180
tccacaacat ccaatcaaaa caacattcaa acagcacaag ctatcacagc caagcaaaac 240
aggacaaaagg cagaaaactc tgctcaacac accaacccaa atcacagctt ttctcactca 300
aagaccccag taacaatttc ttcgatccaa ttcgatccaa gttggatcga ctccaaaatt 360
ttactggaag tctatagtgc ataagcctac attgtgaacc gtggatcta ctaac 415

<210> 857

<211> 591

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 857

agggcacgac ggtnaattga tcgcacatctat tangcgacac tatagatact caagctngta 60
atcatgtgac accctctacc cctcacatgt atactaatat atgaataaaaa ttcaaataatt 120
aattaaaaagt attttaaaa cattttttt tccgaaacaa gtcttcaaa gggaaaaag 180
gctcacattc atttcttct acatcatatt caaacttgc caaataaata ataaagtat 240
ctcgtctcan acaaggtcgt ctaaacttca tacaattaat atagaactta tattctaatg 300
tcacatccctt tcatagtgtt gtgttctgt gtcctctagc atgagggtct tcatagtcat 360
ccacctattc atctgtttcc ccgaacacaa gttcaagatc atcacaggat ccanacacaa 420
caacacacag ggagttagtc atcacattca tagctaatag agagacaaga caattaaata 480
tagatattat ataaatgaga taccacttgc ttaaacatag ctcacgtaat ntccaccatn 540
tgtcattcan naatcacttt tcaatcatca atcacattac acaagaatcc n 591

<210> 858

<211> 416

<212> DNA

<213> Glycine max

<400> 858

gtgtgattcc tttcttttc ttatcattct cctcatgtt attcagtc attagttcca 60
ttcgtgttc ctataactt ccaaataaag ttgcaagaga catgttagaa agatccctg 120
attctgtaat agttgttacc ttgggtgtc attccctact taaacatctt agaactttat 180
taataagatc ctcattggga aatatcttc ctaatgatgc aagatgattt actatgttg 240
tgaatctctt ttgcatatca tgtatagttt cattggatt cattctaaac aattcatatt 300
catgggttaa ggtatttatt ctagaacctt ttacatctat tgccctca tgggttactt 360
gtaaggatc ccacatatct tttgcactct tgcaatttga tactctaaag tattca 416

<210> 859
<211> 487
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 859

catgcgccta tntccttacg aacgttctct ngcacaagac atttagaccg aaaaatgcac 60
ccatatacaa tcaaggcagt ttcggttacct agattattta cacgtacctc caaggtgtat 120
ttgttactta catcacacac attccttgg ctaaattcac atacatgcat actcaaagca 180
ttttggggca ccaaaaattt caccttgca catcttggca tttctaatac ctatacatac 240
gcaaacttca tgatgaatct tgactatcta cacaataagg tgctacattt catgctctt 300
tttcaagttt ttgctaccta aagccgcatg ccaattcaag catattttcc tttgctgact 360
aanatngtat tcaaattaaa aggtatatan cttttgtaa tatagtttct tcacataaca 420
tgcaacatat ttatatatat tttctgtga gacatcttga ctaccaacaa tatataataca 480
tacattc 487

<210> 860
<211> 502
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 860

cgagccncng nnntttagc atctcgaccg cgatccttaa gcacctggc tgcagcttga 60
ccaacagaga gccataaaagt ttgtcttagga agtctaagcc tatagtgcct ggaggcgttg 120

cactctcata ccgtatgcga agagcttcaa gccttgctt ggaagacgcc atagtggatt 180
tcatcctcta ttcttaccc gatcaactatg atagctctgc aggtacgcac gtattgtca 240
tacctatgtt attgttagctc tggcacgacg ttcttcacta catgcggac tgcccaccc 300
agacataggc cacgagtata ccactgtga tcataatggaa ctccctctct ttttaatcg 360
cgatggcatc ccactccgct gctcgttat gacatgataa gacagctgct tcgattcant 420
gtttaccgta atagaagagc acaaccctat tccagcgagg cacatattat tacttgccca 480
ccatgggtct gaattcttgt cn 502

<210> 861
<211> 311
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 861

ttagctctnt cagctacaca tggcggtgac ttgttagagga gtgtgatgga cgaacctgca 60
cccgacgata acactgacaa gagcttatat tctccatctt ggacaagata tggcagactg 120
gtggcaagtc tactatcttc catcagcctt ggaacaactg tgatcgtgat cacatatcg 180
ctaaaacttg atgggtatgc aagccatact ctcactgtgc ttgaatggta acgagcagcc 240
caatgacact gtgacccaca tgtttctcca catgcgttac atcaatgcag tgtctaacgt 300
caagatcaca c 311

<210> 862
<211> 80
<212> DNA
<213> Glycine max
<400> 862

tcctaacgat ttctaattat gtggccatt aagtctatca tatgctgaca atagccgaga 60
agcccatgaa tctcttcggg 80

<210> 863
<211> 440
<212> DNA
<213> Glycine max

<400> 863

cgagaggat gcttcactgg aggagaagac agagggagag atagatagag gcgggagcat 60
gaaattgaac gatgataaac ggagagaagt tgaactatga gttgtgtctc acaagactct 120
cattcatcaa agataacaaca tgtgttacac atgttatctat attatagact atgttagctc 180
cttgagaagc ttcttgaga caactccctt gagaagcttc tatgaaaaa cttcccttgag 240
aagctagagc ttagctacac ataccctct aataactaag cttacctact tgagaagctt 300
ccttgaatag attcctaattg aagctagagc ttatctacac acacccctct aatatctatg 360
ctcacccctt tgagatgaga agctagaact tatctacaca caccctataa tagctaagct 420
tacccatg acacattaca 440

<210> 864

<211> 566

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 864

nttgcgcatttgangcga cactatgata ctcagctggg tatgtcaggc caaatatggg 60
tggagtgaca aaagctntaa ttactgctt gaagtagtgc acgatctgct tccagaggan 120
aacacgttgc ctaaaagcta ctatggcc aagaagatac tatgtccgat gggtatggag 180
tatcagaaga ttcatgctt ccctaattgtat tgcatactgt acagacatga atttgaagaa 240
atgtccaaat gccctagggtg tggngcatca cggtacaagg tgaaggatga taaggagtgc 300
agttctgatg aaaactcana gaagggtcct ccagcgaagg tggatggta tcttcccattc 360
attccaagggt ttaagcatct ttntgctaat gaagacaacg canaagacact tacctggaaat 420
gcanatggga gaaactctga tggatggtc tatcatccgg ctgatntcct ctagtggaaag 480
aagattgatg gttgtattc ggatttcaga aaagaggcaa gaaatcttag gcttggacta 540
gccagtgtatg gaatgaatca tatgg 566

<210> 865

<211> 441

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 865

ataacatcca agaaatttca acatccaaac atcatgaact atcaaaaacta agaaaaaca 60
gggcagaggc agaaaactct gcccaaaaca caaaccaata ccacaacttt tcatttattcaa 120
ataccccaat cacattctt ttgttccaat tcattcaccg ttggatcgac tcaaaaattt 180
tactggaggt ccctagtgaca taattctaaa ttttgaccgt tggttatctcc tagaaaacgt 240
ccagaaccca atctgtacta ctcttccac aaccagcaaa tacacatcat tttctgcatg 300
cacaaagcca aaattctgct gcacattca acagaaaaac tctgcataat agtgcaaaat 360
ttcgaatca cacttgcct tgcctaatt tgccaaatn gaatcctaca agtcctaaat 420
catgtataaa tcatgtctaa a 441

<210> 866

<211> 318

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 866

ataaaagaggg tgaggatgag ggagaaaccc atgctgtgac tgccattcct gtacggccaa 60
gtttccacc aacccaacaa tatcttact cagccaataa caaaccttct tcttacccac 120
cacccagttt tccacaaagg ccatccctaa atctaccaca aagtctgtct accgcacttc 180
caatgacgaa caccacctt agcacaaacc anaaacacca accaagaagt gaatattgca 240
gcgagaaagc ctgtagaatt caccccaatt ccagtgtcct atgctgactt getcccatat 300
ctacttgata attcaatg 318

<210> 867

<211> 471

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 867

agctntgcat ggatgccaca cgtaatctt ctgtatcatc catcatttct ttcattcaca 60
accccaacaa gatgttagtac actagagtta tggcaaaagc aatcagcatc ccaaaaataa 120
cactgcattt ggttaaagaa gaagctccat tggtaggat taaaagagaa agtaattgaa 180

tgaaaaggaa aaaggaatgg aggggagaga aagttctaat tgagccatat aagaattaga 240
tttgaatact cacgctgtgc tgagaatatac aggatgtaca ttatattcct tagcaaagac 300
aaatggaca attccttggg gaagagctgc ctatatttt ttgacattca gttgccaaac 360
aacaggaca agcaaacaaa caaacatgtc aactgcagtt tcaaattcctt ggtgtacaat 420
cacaaacatg atcatngaa tcctcactac tagtactaag atcttcaatt t 471

<210> 868
<211> 338
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 868

ngctngagan actntcttga gaaaacttctt tgagaaactt cattgagaag ctagagctta 60
gttacacacc cctctaataa ctaagctcac ctcccttgaga agttccttga gaaaacttcct 120
tgagaagctt ccttgataaaa cttccttgag aagttcctt gagaatattc ctagagaagt 180
tagagtttat gtacacacac ccctctaata gctaacttca cttcttgag atgagaagct 240
agaagttagc tatacacctc ctataatagc taagttcact cccatgccaa aatacatgag 300
aatacaaaaa cattcctact acanagaact actcaaat 338

<210> 869
<211> 375
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 869

tctatggagg ctggatctt gagttcaat gaggtcctt aatggtgatt ttccaccatg 60
gagatgcagc ggaagacaaa ggaaaagagg tgagaggagg cgccatccac taggaaataa 120
ccatggaaag aaggagctc accaccaaga tgagccttgg ataagaagct tggagaggat 180
gcttcaatga aggaaaagaa agagggagag aaagagggag gggggagcac gaaattgaag 240
gaagaaaaag ggagagaaag agggaggggg gacgacgaaa ttgaaggaag aaaaagggag 300
agaagttgaa ctntgagttg tgtctcacaa gactctcatt catcanagtt acaacaagt 360
ttacacatgc ttcta 375

G C G T T G A T C G T

<210> 870
<211> 457
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 870

ntttagaaaa aatcttataa gtttagtacat acctaaatag ttctttaaga tgactatatt 60
atgcttaaat aatctttaga aatttcaact caataacccc taaagtcatg atattcaatt 120
aaaactatca attattnaa aagagaatgt caacatttgt gatattcaat tgagactttt 180
cacaactaat aaaaaggat tcttctatta aaaaatatat aaattttgat taattatTTT 240
ctagagtcaa ttttggtca attcttagt atgatgtata aattctagac tcatccaaca 300
atttcaccaa aactttcctc atttctgtt aacacatactt aatatgaagt tntgatgatg 360
tcacaagata agcgTTTCTC aagttaatc caagttaaga actcagaaat tcaagataaa 420
tgaagaagta gtccttaaga gtcttagaaa gcattct 457

<210> 871
<211> 417
<212> DNA
<213> Glycine max

<223> unsure, at all n locations
<400> 871

gactaaacat tcattggta tttatttgc ttcattatgc gatataattc gctgtaaccc 60
gtcactaacc aatTAATATT atcaactact cgTTTGGTT agcaaggAAA ttGTTGGTCC 120
aacaaaaatc attacgcgt acagcataca tcattgtcat aattgacaac acataatgac 180
atgcattgcgt gttacagTTT gagcgtgaca acacattggT ngacttcagt acacatTTT 240
aaactAGCAG tcgctcaaca acacattggT tgacttgact acacattgc gacaacacat 300
tggctgactt gactacacat ttacgcgtt ctatttgc tggAAACANAG ttaaacaAG 360
gctcggtcac aaccatctat atatatggca gactangcta ctaaatcaca cattatc 417

<210> 872
<211> 412
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 872

cttgacagct ctatggacca tgctatcatt cttccctgcta gcttagtatt tcttagcacc 60
ttcttgatgg ggtgatcctt cttgacgatg atctaaggc tctggaagta cggttgcagg 120
cgttgagcag aggttatgag tgctagcgcc actttctcga tcatttgta tctcttcca 180
acatcatgaa ggatgtgact gacaaagtag atgggtgtt ggtactttcc atcttcttgg 240
acaagggttg aactaatggc ttttctgcc actgaaaggt ataggaatag ggatgctcca 300
ngcttangtc aacttataac aggtgggtt gcaatagttt tctttatagc tagganagct 360
tgcttacagg cttcgtttca caagaacgac tcggtttcc tgagtagctt at 412

<210> 873
<211> 408
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 873

agcttgcttc tacactaaga agcactctat attgagtcaa tcaccaaaga gagaacaacc 60
accaaaaattg aggaccgttt tgtaatntg taatttacaa tttacttacc ttcatttctt 120
tcaagtttg taacaaaaag gccttcatt ggaagtgtgt tgggagcctc caataagtta 180
ccaaacttcc atttgtgtgt aataattcta ggcaattttt ccttaagata gtgagtgttt 240
tggggaaac cttgaatgtg gtcatccaaa cactcttang attgccttag ttacatttc 300
ttgcttactt tcatalogtta ttcccttac ctccctttt aaaaccacct agatagttt 360
cctttacca attagttttt ttaccttatac ttccacacccctt 408

<210> 874
<211> 321
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 874

tagccctaga ggnatggac cttntcatgt tntggagagg atcaataaca atgcttataag 60
gttggacctc ccataagagt atggagtcag caccactttt aacatttctg atttaattcc 120
ttttgcaggt gaagctgata ttgatgagga ggaactaaca gatttgaggt caaatccttt 180

tcaaggtgga gtggattta atagcacaca aaagtcaagt caatthaact ccttttaat 240
agcaaaacaa gtcaattcta catgtaataa tacaatagaa attgtctcta gctaaattaa 300
aactaaggta atcttgtatc t 321

<210> 875
<211> 463
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 875

gcttgatca aagtgttcga gggggagcaa gcaaagatag aggataactt tcttcttggg 60
aagtttgagc ttttgggtt cactacatcg ccaagaggag ttccacaaat caatgtttta 120
tttgcgttgc acgttgcgttgc catcgtagaa ttcattgcta gagataaaat catgaggatg 180
aaaaaaaaagga tcatgtatcgca caacaagtac tgagggttgc gtccctaaaga gatgaggaga 240
atagttagat atgcaaaagag gtataaggca tangatgtgg aggttaaggc aaacggaagg 300
ccagaacttg cttgagaatt gtgcgttgc aatgtatggc aaagtgaaga atcttaagaa 360
attagtaccc atagcaacaa tgttattnntt tttagttca ttaacaattc agtaaaaaaaa 420
aataccgtgc gctaacttgc aatgccnct gcacacatgg ata 463

<210> 876
<211> 510
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 876

atccttatgg cctgcctnca gacttcaccc ncgcgtgcca ccccgaaaga attaagccaa 60
gcccctactt ttgacggaa actcccacct tatgaagact atcccgggca agacgtggg 120
gagggagata cccatcttgg cccctgctc cacctcaaag atccatcccc acatgaacta 180
ccccagccga acatagtccg ccatatcccc gtctcacccca cacccgtaaa agaatctgtt 240
cccttcgcgg aagataaggg aaagattgag gcgcgttgcg agaggttaag agcagtcgag 300
ggcctcggtt attaccatt ctcgatattt gcagaattat gtcttgcgc caacattgtc 360
atcccttcca attcaaagta ccaaactttt attagtagca agggatgaca tgttcaaang 420

ggcatctcg atgtatgc tgagatggng catattctgc ggacaannag tcgtggcat 480
ttctttcag acaggttgct tggacngctg 510

<210> 877
<211> 383
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 877

agcttagccc tagagggat ggacctttc atgtttgga gaggatcaat aacaatgcct 60
ataagttggg cctcctagaa gagtatgcag tttagcaccac ttttaacatt tctgattaa 120
ttactttcc aggtggagct aatattgagg aggaggaact aacaaatttgc agatcaaatc 180
ctcttcaagg gggagggat gatgcaatcc tccctaggaa gggccagtca cttagagacat 240
gagcaagagg ctccaagagg attgggctag agctggtgaa gaaggcccta nggttctcat 300
gagcctcatg gttagattct gagccatgg gacaaggttg ggtctaatta tctttgtaca 360
tattaaacta ngatgtcatt ata 383

<210> 878
<211> 490
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 878

ngttgtgtt gtggacctct atnntaaaaa cctccataag agttgttaga aggacctaca 60
aaagaagtgt tagtgggacc cacatattgt ggttccagc cataaggctg atttgatctt 120
gagccaccc tcataccctt gtagcttcct tgaaagttnt gttgaggtct agcttggttc 180
tgttagatatt cggttcctc ttcttggc catcaccagg tgctgaacag tggccattct 240
gatggataacc accacaaaaa tcacatctca naattntttg aacttggtg gcttgatatg 300
ttttctatgg tccacccctcg tgatattgtc gagtcagttg gcctatctac tttgttaagg 360
gcctcaatttgc ttgcgtcaag agnttggtnntt gagctagaat ntcactctga gtgtccagct 420
ccattataacc tattcttgc gtcggatcnt catcatgtat actttgatata cactagctac 480
tatggagttt 490

<210> 879
<211> 472
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 879

agctatgtgg caagtacaaa ggtcataaac gcgcggta tattatcagg tcttgctttg 60
gtggccttaa gcaagctttt attgctagtga gaagtggat ttcacaggta tgaatcttca 120
atcatcattg atcaaaatgg gaaaatattc gccagtgtaa tttttataa tttgcaaatt 180
gaaagccttg atgttatat gtctctgtta aatgtgtttt atttgctaag tttttatagc 240
tgtctcaata atttggtaaa ataagttcaa catgcacttg atgcattgcta tcgaggatca 300
ctaaaaatatt ggcataaaag acccatgaaa tggttcttg tggtctgatn tactggactt 360
gaatgaattt aactacacat cgctataatg ttcaagagtt cctggcttct gcaatattat 420
.tctagtttat cttgataaaa ctaggaacat ctgcattgtat aatgctggaa gt 472

<210> 880
<211> 284
<212> DNA
<213> Glycine max

<400> 880

cagcttctcc attatctatg ttctcgattt tatctagcaa ccaagttagg gtggagttct 60
tattatgatc ttgtaaagct gagccattgg ataccaattc atccttagct tgcgaggata 120
agtgcgttaa tcctcccatt agaatatgca tatgcgtat cgctgtatct ctatcaacaa 180
attcgtctct gtcttcgtat tccatgggtt ccatcaacat accatcaaacc atctcgatct 240
ggcataaaatt gtctatcatc ttttgcgtat cctgcctagg atct 284

<210> 881
<211> 410
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 881

tattccatat tgggaatct aaaataacag acaattgatt gtacaaaaca atgataggag 60

taaaatctag taacatcata atttaaactc actgcaacgg ataaaatgaca atcaccatac 120
tcaagagcct catggataat gataagatct tcactaccct aatctaagaa aatttagacga 180
taaaacaatn tatgttttgtt gattacatag tggcactaat atgtaaaatgc aaaaatgctg 240
acctggccat gttcattgaa agtatcgagt ccaacaattc caaggtaag atctccagat 300
aacaattttc ttgtgatgta ttgggcctc taacaccaa ctttgagttt ggatagctgc 360
aataatgcaa gagaatttgtt ttatggcat gttagagatat agtacattga 410

| | |
|-------|-------------|
| <210> | 882 |
| <211> | 371 |
| <212> | DNA |
| <213> | Glycine max |

<223> unsure at all n locations
<400> 882

tcccgatct gatcatggaa ggactnggca actgccttca ttatgcagta ccagtacaat 60
acggacatgg ctcccgatcg gaaccagctt cagggtatga ctaaacgaga gcatgagtcc 120
attaaaggaaat atgccc aaag atggagagat tntgcagctc aagtcgtacc gcccatgatg 180
gagagggaga tgatcacaat tatggtagat acgttaccca cattccacta taaaaagctg 240
ataggctaca tgccagctaa ctttgcggat ctcgtttcg ccggcgaaag gattgaatcc 300
gggctacgaa naggcaagtt cgaatatgct gccaatatgg ccccaacaa caagagaaga 360
cccccaactg t 371

| | |
|-------|-------------|
| <210> | 883 |
| <211> | 254 |
| <212> | DNA |
| <213> | Glycine max |
| <400> | 883 |

| | | | | | | |
|-------------|------------|------------|-------------|------------|------------|-----|
| <400> | 883 | | | | | |
| cgcgaacttt | gaccattgtt | cttccttccc | gcaatgcttc | ttttcatgtc | tgcctgagtg | 60 |
| ggcttatagc | ctaaaccata | cttcccacga | tttcctttag | tatttatcag | gctagttatg | 120 |
| ccgcccgttgt | ttttcctaa | accatcccc | ggttcaaaaac | cgttccccaa | cataactcgg | 180 |
| gccatcatta | ccgctgcata | ggacagacaa | ggttgcccaa | agaggagtc | cacggaggaa | 240 |
| atgctgacca | cctc | | | | | 254 |

<210> 884
<211> 101
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 884

ntgaggggtgc gtagccacc atctnttcat agtagagtgat cgataatgtg tctaccatca 60
cgattatcg ttcctttcc atcattggg gtaccacttg g 101

<210> 885
<211> 300
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 885

tttccatatg tcataaca aaaaaaggaa aaagggatg tagtggtga tgcaactgtct 60
aggagacatg ctttacttgc tatgcttcaa actaagttgg ttgggttcga gtctttgaaa 120
gacatgtatg tgcatgtatg ggactttgct gaaatttttgc ctgcattgtga aaagttttct 180
gaaaatggtt actataggca taatggattc ttggtaaag caaataaatt gtgtgtgcct 240
aagtgttcca ttagagagtt gcttggatg gaatcacatg aggnggggtt gatggacac 300

<210> 886
<211> 331
<212> DNA
<213> Glycine max

<400> 886

tccattgtcg aatttcgagc gtctcgatat atgatgcgcc taattctgac ttccgagtga 60
agagttatga ccatttgaat tactggcgag ctccgttga tcaatttcga gcatctccaa 120
acattatgcg ccttaatcgg acatccgagt gaaaagttat gaccatttga agttctcgag 180
agcttccgtt gttcaatttc gagagtctcg atatattatg tccgtgaatc tgacattcat 240
gagaaaagtt atgaccactt gaatactcga gagcttcgt tgtcgcattt cgagcgcctc 300
cgtatattat tcgcattaaat cggactttct a 331

<210> 887

<211> 233
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 887

aactggggca aataaagagg gtgaggatga gggagaaacc catgctgtga cggccattcc 60
tatacgcca atttcccac caaacccaac aatgtcatta ctcagtcaat aacaaaccac 120
ctccttaccc accacccagt tatccacaaa ggccatccct aaatcaacca caaagcctgt 180
ctaccgcact tccaatgacg aagaccacct ttagcacaaa ccananaaaa cac 233

<210> 888
<211> 336
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 888

tccattcccg agagcatntc ttatttaagc acnttcagcc ttgcttcat gtagcttagg 60
aaaaacatca ttttttttcc ttttttttcc ccaaagccaa ttctaaagtt ccaagcactt 120
tctccatcac ccacagccac cattagccac cacaaccat cggtttctc cattgaaacc 180
ccacaccgag aggaaccctt caaccgaagt ggaatcttcc aacttggctt gcggtttcgg 240
tagagaacaa aaccctaatac tgacccttcg ttttcttttg agactatntt agtctcaaaa 300
ttatcaagaa ctacgttaggt ctgagttcct catcac 336

<210> 889
<211> 563
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 889

attgaaccca tttaangccc tctattagct gaactctcat acatacgtct gagccggact 60
ctgactctgc attaccttgt cccangtgat aatgtcaatc cttactcttc gaagcancga 120
ggaaacgaga gaaggataat ttccactcta aggacataag gagagggaaag gatattcctc 180
atcaaagagt gggagatagc tatacgacca gatagataat tcccaatcca agactgtgag 240
agagaacaag agaccgagat gacngaagga tagctcctga tcaatgatcg aaagataaca 300

gaagaaatgt gcagagggga tctctggaca gacaatatct atacaaatac agaattgtca 360
ccaaatgaac acaagagaga aaggaaacca taacctacaa gtggtcttct gccttcgatt 420
accaacccaa atactgtgcg tcngtgactt ttgtcgctcg cgtcagacaa naactgaaaa 480
cgaaaacagc cacactaaaa ctatcaaaag ccataacaac aanagccgat aacccactaa 540
agagtcatcg cacgggagtc tat 563

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<210>      890
<211>      471
<212>      DNA
<213>      Glycine max

<223>      unsure at all n locations
<400>      890

agttccatc anatggtata aaaggcacaag agcttaaagt acgtgctcct taaacacctca 60
ttaaccttta ttgggtttc ttcattttc tccatgtata tcctcacgtg tcttgctcta 120
aatgttgttc acatgatttt ttagaatttc gacaattaaa cttgctatag aagctagatt 180
tgattttcta tggttcaaattt ttcttggttct tggcttgaa ccataattgt gttgactnta 240
ngtttctttg agttttgtct tgctatattat ttgtggctga aacttaaacc ataaaattct 300
tacaaaaata ttaatgtata agaaaaacctc aaaaatctag agtgacatgt tcacctattg 360
tagttntgtc ataaaaagtca tgtctagtca tggaaacttgc catatatgtat tctttatgtn 420
ggacttggaaattt ctcattttct tgggttcttgc tctaactcat ttgtccttgc g 471
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<210>      891
<211>      461
<212>      DNA
<213>      Glycine max

<223>      unsure at all n locations
<400>      891

tcaacatcg accacttcca gggtgctgga actacttcac atggacttga tggggcctat 60
gcaagttgaa agccttggag gaaagaggta tgccatgtt gttgtggatg atttctccag 120
atttacctga gtcaacttta tcagagagaa atcagacacc tttgaagtat tcaaggagtt 180
gagtctaaga cttcaaagag aaaaagactg tgtcatcaag agaatcatga gtgaccatgg 240
cagagagttt gataacagca ggtttactga attctgcaca tctgaaggca tcactcatga 300
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gttctctaca gccattacac cacaacaaaa tggcatagtt gaaaggaca acaggactnt 360
gcaagaagct gctanggtca tgcttcatgc caaagaacct ccctataatc tctgggctga 420
agccatgaac acagcatgct tcatccacaa cagagtcaca c 461

<210> 892
<211> 465
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 892

agcttgacca atcccgaccc aaccgggca tagtcggtca gtgagaacct gtgatgtacc 60
taagcaggcg agtcctggc agtcaacaga taaaaggaac aaagaccaca aagcaaggag 120
gcttgtggc gctggccagc tgtgaatctt gtgtgatata tggtttttgc cctctggtaa 180
tcgattacca agggtggta atcaattaca aggctaaaa atgaagacag gaggctaaga 240
tggctctgg taatcgatta ccaaagggt gtaatcgatt accaggcttgc aaaacgagg 300
caggaggcta tgagggcttc tggtaatcga ttaccaaggg ggtgtaatcg attaccaggc 360
ttaaaaatga angcagcang ttgttagaggc ctctggtaat tgattaccag tctgtgtaat 420
cgattacaca gaggaatggg tcactggtaa tcgattacca cgtat 465

<210> 893
<211> 238
<212> DNA
<213> Glycine max

<400> 893

tttgggctag cccatgttcg atactctaca tagaggttagc gtggAACATA ctttgcAACAA 60
gtgtgtatac ataggtaaat ataatgagca tggaaattcct agtaaagtgt gaatgattgt 120
cttcctaaat gaatgttatga tagtgtggaa tgccttttgc aatgcAAATA tgtgcATGAT 180
gtaaatagct atccaatatg catataaata aatatgagtg aaacaataac aatttgt 238

<210> 894
<211> 419
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 894

agcttgttccctctaaggta cttgattctt ggaaagtgtatgtccatcat tttcttctat 60
tttcttagacc cttttgcac cattttaatt actgattggc cttaatggc aattaatcag 120
gcagtttat tatgggct cattagcta atttgatgtt tttaatctaa ttttaggaat 180
taatgaaaca ttgggcttaa tccggatttt ggatatggac ttgaagaggg caaataaagc 240
agcgcttatac ttagttaatt tctaattagg aaatttgca attttatccc atgttggca 300
gtgttattt cgtnntggc cagagtattt taatagagcc cagtgacttt gagtgactct 360
ttntaaatag cagccttggg attcgtgcaa ggcattctat tatgctatcc tcattattc 419

<210> 895
<211> 402
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 895

tatccncata agagtgcaga acagctggcg agtcagcatt gattatagga ggctaatcca 60
ggtaaccaaa anagatcatt ttccccgtcc attcattgtat canatgcttgc agcgcttggc 120
aagtatgtct cattacaatn ntttatggt ntttctgggtt atttacaaat tcataattgct 180
cctgaggatc aagaaaacac cacattcacc tatcccttgc gcattttgc ctataggagg 240
atgccctntg gcctatgcaa cgcctctggt accttccaac ggtgtatgct tagcattntc 300
aatgattntt tagagagttt catagatgtt ntatggatg attntactgt ttatggatcc 360
tcttnngat gcatgttggatgtctagat agagttctta at 402

<210> 896
<211> 396
<212> DNA
<213> Glycine max

<400> 896

atcctctgag tcacctgcgg ctgcagcttc tatccaggct catcttggtg gtgaagctcc 60
ttcttccatg gcttattccc tagtggatgg cgcctcttct cacctcttct cctttgtctt 120
ccgctgcatac tccatgggg aaaatcacca ttaaaggacc tcattgaagc tcaaagatcc 180

agcctccata gaagccccac aagcaagctt ccatcagagg aggagtcac ccctctttag 240
ccttcttatt ttgatgcagg tacgcacatg gctaaggagg aggataacctc cacaaccag 300
attcatgagc catcttctac acttgtacct gatgatgcca caccatctgc accagcacct 360
gagtcagagc atccttatctc ttaagattca ccaact 396

<210> 897
<211> 354
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 897

tgtcatgacc acgctctctg gtgcataatt gatctgtcca aacttaacag ctcattctnt 60
tcgaacatac tgaataatca gccaggagct tacaagcgta caagtggaaa aattactcaa 120
ctcttacagt atgttctaag tctgagtaat ggaatacat attgcttact atttaactac 180
aacgcttact tgacagatga gcagttcatc atagctatcg gaaggtgaca ttatcttta 240
ttcttaatta ccccttaatt tgtacatgca ttattaaaca acctttaaa acaaaaatac 300
ttcatcaata ttagctctca agtctaaatt agatgccatg tatcatattt atat 354

<210> 898
<211> 446
<212> DNA
<213> Glycine max
<400> 898

agcttatgct gcaaacatct acaatagacc ttctcaacct caacaacaaa atcaggcaca 60
acagaataac tatgacctct ccagcaacag gtacaatccc ggatggagga atcatcccaa 120
ccttagatgg tcgaatccgt cacaacaaca accttatttt caaatgttg ttggcccaag 180
cagaccatat gttcctccac cattccagca acaacaacaa caacaacatc cccagaaaca 240
gcaaacagtt gaggccccctc cgcaacccctc ctttgaagaa ctttgtgaggg aaatgactat 300
gcaaaacatg cagtttaac aagagaccag agcctccatt cagagcttaa ctaatcagat 360
gggatagttg gctacacagt taaatcaaca acagtcccag aattctgata gataccttct 420
aatctgtcag aatccaaaaa tgtgag 446

<210> 899
<211> 360
<212> DNA
<213> Glycine max

<400> 899

tgtccgcaaa agatcactaa caacgattct aatgttcgag acctaattt tcttcacca 60
agtaaaaaatg gatcattcta aggtccaacg cttataatg aacacccccc aagtaaaaaa 120
aatagcttga ttccccctta aaaagaacta cgtatgtctg atttccttctt cgatggagg 180
tacgtagaag caagagccct gctttgtcg acctcacaaa taaaaaagaa ataaaaagtt 240
tatgtacaca atttcataca attcaataat taaggctgtt gtccttgag acaaacgtga 300
gaggtgctaa tacccctc aaacgtaaat acaactcccg aatctggaat attcttcatg 360

<210> 900
<211> 449
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 900

agctnttaac cttgcttctg tgtatgcaat tttaacaggg gacatatttt atgatgctag 60
tgaagatttt tctggactg tatctgttga tgatggtgat gataatatct gcagaatttt 120
gactgtgaa gatagttgg gtaccaatgt tggagtataat actgatgata cagaagaaac 180
aacagacatg ctacatgcac ctcctccccc tggaccgaat aagagaagaa aattaatgaa 240
ttccttagt gctggagttg aagttgatag ctactcgaca gctgaaattt ataactcatt 300
ggattattct cagacctcta gctgtgttcc tgatgataca gttgaaacca ctcaagatga 360
tacactngaa accactcaag atgatatagt tgaaactact caagatgata cagttgaagc 420
aacacaatat agtcatggc tactgtcat 449

<210> 901
<211> 399
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 901

ntgagggtgc gtagccacc atctttcat agtagagtat cgataatgtg tctaccatca 60

cgattatcggt ctccctttcc atcattgngg gtaccacttg ngccgcaga tccctcaacc 120
tttgggcgt gttcttgaa agatccgtcc cccttntgc aaatgttctg tagttgcac 180
ctatccagaa ccatatcaa attgtactaa tactgcctaa caaaggcaac caataggc 240
ttccaagaat ggactcggga aggttccaag tttagtgtacc gggtaacagc tacccagta 300
agactttctt ggaaggaatg tattagcaat tcctcatctt ttgcgtattc ccccatcttc 360
tgacaataca tcttagatg gttcttggga caagtagtc 399

<210> 902
<211> 565
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 902

attgacgcat cttangcat cactatcgat actcaagctn gcctanagag gtccaggana 60
gacaaggcgg ccgaaggaac tagttccgcc ccgtagtacg acagtacccg cttaggagc 120
gttgcacacc agcagcgaaa cgaagccatc aaggatggt cgtttctccg agagcgacgc 180
gtncagctca nggaggacga gtatanctga ttccaggagg aaatagggcg cccgcgggtgg 240
gcaccactgg ttactccat ggccaagttt gatccagaaa tagtccttga gttntacgcc 300
aatgcttggc caacagagga gggcgtgcgt gacatgagat cctgngttag gggtcagtgg 360
atcccgttcg atgccgacgc tatcagccag ctccctggat atccgatggt attggaagag 420
ggccaggaat gcgagtatgg ccagaggagg aaccggcttg atgggttcga tgaggaggcc 480
atcgcccagc tgctatgtat anccngntan ngatttgcen gactgctgan ggagagagtg 540
cgatcatcnc accatatgac acccg 565

<210> 903
<211> 381
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 903

taggggggag aagtgaagaa gaaaagggtt cagcctctaa ggcacttctc tctttctcga 60
aattgcttag gaaaattatt ttcgtgaaga aaatccaagc cgaggcgctt ccgtaacgtt 120

tctgttaacgt ttccatgagt aattacgcga agattctcgta ccgttcttca agattcatcg 180
ttcgttctgc gttttcttca gtcttcaacg ggtaagtacc tcaaaccagg cttttcaatt 240
cattctatgt acccggtggt gtccacattt tgtttcatgt attttatttc tcattttcat 300
ttactttnta tacccccc ttgacgtgctt aagccattta tttaagtcat ttctcgctta 360
ataaaaaat aaaataaaatc t 381

<210> 904
<211> 387
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 904

cccatcacat gtggtactat gtgggtgtcg ggcgtatggtg cataatgatt ctccacatcc 60
acaatcacg tataacccac catccccctgg tgcccacctc caactgagct cacgtactcc 120
cacgttagccc ttatcctcgn tcctctcaac gccgggtccc catcaatcct cccaagcttc 180
cacaacatcc aagtaattca acattcaatc atcacaaact aacacagcca agaaaatagg 240
gccaaggcag aaaactctgg ccaaaacaca aaccaacatc acagctttc acattcaatt 300
acctcaataa gagtctctgt gttccaggcgt ggtaaccgct ggatcgaact cgaaattata 360
ctgggaagct cttagtacata agtctac 387

<210> 905
<211> 130
<212> DNA
<213> Glycine max

<400> 905

gagcctgtt tcccttcct tgtttgaag ctcactacaa gccttaaatg aaaaaccatg 60
atatcaccat atccttaagg aattttggag ctttggatt gttttggaa taagtgtggg 120
gggggggggg 130

<210> 906
<211> 417
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 906

tattacggac ctatagatac tcagcttcaa gacaaccttg aaaagatggtaattagcttt 60
ntccttcaca ttcagcanat tcagcattta aatgtgatat ttaatgttat gctntttat 120
gccatgataa ttggtgaaat gaatattatt tatttgc当地 ggttcatga tatcgaatat 180
tgatacctaa naaggtaat attcaagtt gtgtgattag tgttattttg agatgaaaca 240
ccaactatac gtaatcttat cttgcatta tcaagtttgtt attaaaaatt tgtaatctat 300
tcgttggata tgatagtagt agggactcat aaggatntac ttagtaagag gcttaaccta 360
aagtaagaat ttgttttct gagacaaaac tgcaagatc atcntgtttt attatta 417

<210> 907
<211> 465
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 907

cgtgcattca atatcctgat gaggaggttc catatgtctc aagactggac taatacattt 60
gctgtccaag tttcatggtc ttgcaggta agatcctcat aagcatctta agaagttcca 120
tattgtctgt tccaccatga agccccctga tgtccacgaa gatcatatct ttctaaaggc 180
tttcctcat tctctggagg gagtggaaa agattggctg tactaccttg ctcccaggc 240
cattaccagg tgggatgacc ttaagagggt gttctgggg aaattcttcc ctacatctag 300
gaccactgcc atcaganaag acatttcagg catcangcaa cttagtggag agagcttgc 360
tgagtattgn gaaagattca agaaattgtg tgcaagttgt ccccaccacc agaattttga 420
gcaactcttt ctgcaatatt tctatgangg acttancaac atgga 465

<210> 908
<211> 588
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 908

gggaccacgg gnnnnnaatt gatcgcatgc tattangcca tactatagat tactcaagct 60
cgagttgagg aagtgttagaa gggtgaaact tcctggcttt attcggtgac cacagagtgg 120

tacctggaga tatgtcgca gggtcaggag aaccttggga cgtcaggtgg tgtgctattg 180
cccaaaacca agcttgacca atcccaccc aaccggggca tagtcggtca gtgagaacct 240
gtgatgtacc taaacaggcg agtcctggc agtcaacaga taaaaggaac aaagaccaca 300
tagcaaggag gcttgtggtg gctggccagt tgtgaacttt cattgatatg tgggttatgg 360
cctctgttaa tcgattacca agggtggta atcgattaca aggctaaaa atgaagacaa 420
gaggctaaga tggtctctgn gtaatcgata ccacggngtg taatcgatta ccaggcttga 480
naacgaggc aggaagccat gagggcgtct ggttaatcga taccaagggg tgtaatcgat 540
taccaggctt ananaggggg atggacattt tganggctct gtaatcan 588

<210> 909

<211> 267

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 909

agtatgacag tcaccgctt aggagcgtt tacaccagca gcgcttcgag gccatcaagg 60
gatggtcgtt tctccggag cgacgcgtcc agtcaggga cgacgagttat actgatttcc 120
aggagaaaaat aaggcgccga cggtggcat cactggttac tcccatggcc aagtttgc 180
cagaaatagt ctttgagttt tatgccaatg cttggccaac agaggaggc gtgcgtgaca 240
tgagatccta ngtaaggggt cagtgg 267

<210> 910

<211> 361

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 910

tacggatgga atacttactt ggtgtgatga acaagagcgc gatacagaat ctataaatgt 60
gcaaaatgat gaccctangg ctgctaactc gtaaatcccg tgggtatggc tcttggaaagg 120
cgaaaaaaga agtttatgaa tgcaaaaacg cgccccctt cgtcattttt atatatttgtt 180
gcangggtgg ctcgcccagg cgagctaacc tgcattttt tttttgagag gaacattaac 240
catgtccact cttccctttt agcgcttgc ctaactgaa cttacttaag ttägaatcaa 300

gcgttgatta cttatTTTA ataacaaaca gatAGTAAGA taACTGCGAA tacAAAGGAT 360

a 361

<210> 911

<211> 471

<212> DNA

<213> Glycine max

<400> 911

agcttctcct actgcaattg tcaaACACGG tgtCTAAATG tgAGTTCGAA taACAACCTC 60

ttttACATGG tGATCGCGGA tGTACATCCC aACTTTAGTC atATTCAAGAT tCTCGTTCAA 120

actAGCGTCC accATTGCAC tCCAACCTAC tCAATATAgg aggGCTCCAT attTCATTG 180

tGCTATAACC acGCATTCTC tCTAAATCTC CCCTGCACCT tTTTCATTG aACCAACCT 240

agaaaaAAATA ttTGCATAAT gaATCACCTC CGAAATCTGTA CTATCCTTAT tGCTCCAAAG 300

cTTTCATTc catCTCCTCC aaACACTCCA taACATCATA gCAACACGTT tCCCTTGCCG 360

gACTGATAGG ACTTGAATTa ATCTGAATAT gaATTCAAAG cacGAACTCT CATCAAATAT 420

cGTTAGGTCC tCAACAAACAT CCCAAGTGGa agACTCTAA gTTAAATATT T 471

<210> 912

<211> 409

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 912

tATGCATTGT ggaATTCAC cAGAGAGAGC ATTGTTGAA ACATCTAAC tTGTGATGC 60

tTTCAGCTTA ACCGACTTCa aAGGGTAGGC tTTCATTGA ACAGGTGTG ggACAAGTTC 120

aATAACAATG ACAAGTANGA aAGACCAATG ACTTGCAAAG GAATAGTTCC ACTAATTCTG 180

ttGTGTAAA gGTCAAGGGT ATTGTAATT TGCAATTCC CAACACCAGG ATGTATGCTT 240

cCTTCCAACA CATTATTGA CAAGTCAAGT tGAAACAAAA GACTAAGGTT GCCAATGGAT 300

aatGGAATTt CTCCTGACAG tTTGCTCACA tTTAAATTCA ATGACTGCAt CTtTTGGAAc 360

atGACAAAG AAGCAGGAAT AGTCCCAGTA ATCTGATGGC CACCAATAT 409

<210> 913

<211> 442
<212> DNA
<213> Glycine max

<400> 913

gatcttaagc acctgcggct gcagcttgc ttaagacatt gtctgggtt tttgcttctt 60
tattttttc tgaaaattgc tagtttagta taggtccttg atttttggtt tatttgaat 120
aatgtgtac tccttgtt tgaggctta agcttaagta tagagtagtt gcttcaaga 180
atagtgtgc tatggaaatt tccttaaat ttgcggcaa cgtcaaacc aaatccctacc 240
caatgttttgc aaatccatca tactgcgcct ttagaattcg aagaatggta caatgatttt 300
aatgggtccc accactgggt tatactgtat taaatatcca ataaatatac ataaataatg 360
gaatcacgtc ttagacagaa atgttacata acaatactac aaataatcac atactactat 420
gctacgaata atcactgatc tg 442

<210> 914
<211> 524
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 914

ngagatgagg aagtgttcaa gggtaact ttctgtntt attgttgcacc acagagtgg 60
acctggagat atgtcgccgn ggtcaggaga accttggac gtcagggtgn gtgctattgc 120
ccaaaaccaa gcttgaccaa tcccaaccca acccggcat agtcggtcag tgagaacctg 180
tgcgtaccc aaacaggcga gtcctggca gtcaacagat aaaaggaaca aagaccacaa 240
agcanagagg ctgtgggtgg ctggccagct gtgaactntg attgatatgt gggttatggc 300
ctctggtaat caattaccaa gggtggcaa tcgattacaa ggcttataaa tgaagacagg 360
aggctaagat ggtctctgg aatcgattac cacggngtgt aatcgaatac caggcttcaa 420
nacgagggtca ggaagctaag gaagcctttt gtaatcgatt accaagggtt gtaatcgatt 480
accaggctta naaaggaaac tggagatga tggaaaggctc tgg 524

<210> 915
<211> 305
<212> DNA
<213> Glycine max

<400> 915
ggctgcagct tattgttaca agccacttac tcttttagcc ccaagagact cagcataagg 60
atgcacagac caaaaggtttg tttgtaaaaa aatgtgttga ccaatggaag gtgctaattg 120
caaaaacaaa tgaaagctat gccaaggcaag cccaaaaaaaaa aaggaaggaa gtggttcttg 180
aacccggaga tggatcttggg cattcgagga caaatgtttt ccaaggaa gggaaatgttg 240
agaatcatga aacaggccaa atacagtctt aaggcccag tggagaagga cgaaggcccc 300
agtgg 305

<210> 916
<211> 353
<212> DNA
<213> Glycine max
<400> 916
tcattgccta acaaggccaa ac ttacaacatc tagccccaa agactcatca taatgtatc 60
caggtcaaag ttgagttatga gataagattt tatgaccaaa tgaagggtca tattgcaaag 120
aacaatgata gctatgccta gcaaggccaa ac aagataagga atgaagtggt acttgaaccc 180
tgtgtatgtc ctggacattt gaggacaaat gttttccaaag aaggaggaa tgatgagaat 240
catgaaactg gccaaatata cgctaaaggc ccaagggtgg aaagactaat gcctgagtgg 300
agaatgacaa taaccctgag tggagaatga tggaaaggccaa agtggagaat gat 353

<210> 917
<211> 404
<212> DNA
<213> Glycine max
<400> 917
actcagcttgc tcatgaccccg tctctttggc gcattatttttga tctgtccaaa cttaacagct 60
cattctttttt gaacatactg aataatcagc caggagcttca caagcgtaca agtggaaaaaa 120
ttactcaact cttaaaagtat gttctaaatgc ttagtaatgg aaatacatat tgcttagtat 180
ttaactacaa tgtttacttg acggatgagc agttcatcat agctattgaa agtaacattt 240
ttctcttattt cttaattacc ccttaatttg tacatgcattt attaaacaac ctttataac 300
aaaaatactt catcaatattt agttctcaag tctacattaa atgccatgtttaaaatattt 360

ataaaaagttg ttttcatatg ggattgataa gcgtgtgtgt gtct 404

<210> 918

<211> 453

<212> DNA

<213> Glycine max

<400> 918

agcttggcg caaccaccc ccttttttc tctataatag gggaaaaagg gcagagtaat 60

ttggctcaac ccttctggaa tttaggattc tcttgaattt agagagaaaa attgtttccg 120

tgaagaaaaat caataccgac gccctccgt aatgcttctg agacatttc gtgagcgatt 180

ttgtaaagat tcttcaccgt tcttcatcgc tcttcgttcg ttcttcgtcg ctcttcggc 240

ttcaaccggta aagttccctga aataaaaccttcaattcat tctatgtgcc catagtggc 300

cccacctgtt tcacgtgctt ttatttcat ttctgttgc ttccgtaccc cttttgacg 360

tgctttaacc attatTTAAG tcgctttctc acctaatacaa gtaataaaat gagattccac 420

caatcatttgc agttgtataa tcgtttaatc tct 453

<210> 919

<211> 419

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 919

gcngcaccgc tnaaggtaaa gactcttcgttca aacttccaaat atggtcattc 60

aggccctcac cattctgctc cttcttgaa tatggcaat ctctctaaat atgcccctct 120

tgtcaatagt tgatacaagt cgcccttta tcagcataat tcgagggaaat gtgcctggc 180

ttaccacatt tgtaacaagt gatctgagtt gataaagaag tgggttgct accattacca 240

ccagcaaacc ccatagcatc agtcctctga ttgttggggc gattaccata tgtcttaaga 300

ggggttgagt acgatcttcc ccgttggta ggtccattct ttttgttctt cattngcct 360

gcactcctat aatacgctgc cttgtatcag aagcttcatc ccaatccgga catgttacc 419

<210> 920

<211> 245

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 920

cacaacaagg tttcacatcc acaatgcgac cataaaaccca ccatccccctg gtgcccaccc 60

ccaaactgagc tcacgtactc ccacgtagcc catatcctcg tttctctcaa caccgggtcc 120

ccatcaatcc tctcaagctt ccacaacatc caagaaaaac aacattcana cagcacaagg 180

tatcacagcc aaagcaaaaaca gagcaaaggc agaaaaactct gctcaacacaca tcaaccagaa 240

tcaca 245

<210> 921

<211> 397

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 921

cgcctaatta acctgatatt gagaganaat gattattaaa tacacaaaat ggaagtacta 60

agtattttt atcttatattt aatagaaaaat acttataaca ttacaaaata accataaatt 120

ggaagagttt gatacaattt acacaagttt tatacacaaa agttcgctgt attcaccgac 180

taacatagca caagacatat ccgtggaggg tttcgagggt atagtcaata acatcaccgac 240

caacaattac ctcactttcg ctgacaaaaga gatactcgac gagggcaggg gacacaatgc 300

acgtgtctgt caaatgtttg gacaacatan gggccaaagt gctcatcgac aatggctctt 360

ccctcaatgt catgcncaa a gctactttgg acaagct 397

<210> 922

<211> 406

<212> DNA

<213> Glycine max

<400> 922

ggctgcagct tctcgatata ttatgcgcc a gaatcgacc tcagtgtat aagttatgac 60

cattttgaat ttccatgtt caatttcaag cttctcgata aattatacgt 120

ctgaatcgaa ctttcgtgt ataagttatg accatttggaa ttcctcgaga gcttccattt 180

ttcaatttca aacttctcgaa tatattatac gtctgaatcg gactttcgat tgataagtta 240

tgaccatttg aatttctcg a g g c a t c c a t t g t t a a t t t c a a g c t t c c c g a t a t a t t a t t 300
g c a c a t g c a t c a g a c t a c t g t g a a a t g t a t g a c c a t t t a a t t t c t c g a g a g c t t c c 360
g t t g t t c a a t t t c g a g c g t c t c g a t a t a t t a t g c g c c t g a a t c g g a 406

<210> 923
<211> 393
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 923

c t a t t a c g g a c a c t a t a g a t a c t c a g c t a g a t t g a a c a c g g a a g c t c t c a a g a a a t t c a 60
n a t t g t c t t a t a c t t c a c a c g g a a c a c c g a t t c a a g c t c a t a a t a t a t c g a g a c t c t c g 120
a a a t t g a a c a c g a a a g c t c t c g a g a a t t c a a t g g t g a a a c t t t c a g a c g a a a g t c 180
g g a t t c a g a c g c a t a a t a t a t c g a g a a g c t t g a a t t g a t c a a c g g a a g c t c t c g a g a a a a 240
t t c a a t g g t c a t a a c t t g t c a c a c g g a a g t c c g a t t c a g c t c a t a t a t a t c a t a g a g a c g 300
c t g g a a a t t g a a c a a c g a a a g c t c t c g a c a a t t c a a a t g t c a t a a c t a t t c a c a c g g a 360
a g t c t g a t t c a g g c g c a t a c t a t a t c g a g a c t c 393

<210> 924
<211> 382
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 924

a t g c c c c a c a t t t c c a t g a c a c a a a t g c a a t t g g a a a a c t t c a t g c a a 60
a a c t t g t c a t g c a t c t a t g c g g a c a c t c a a t t t t a t g g t c a t g t g a t 120
g c t a a g g c t c a g g a t t c t c t a t t t a t c a a c c a a t g t t c c a a a a t a t g t t 180
c t t t t a t c a a t t g t g c a t t c a t c c g a g t c a t t c g g g c g t c c g g g a a a t t c a c a g c a 240
t t c a c c c t c a g g c g t a g a c a c a t t t c c a a a a t t g g t t a t g g t c a a t g a t t t c a 300
a a g a a a a g t t g g a a a t c g t c t c t t c a a a g c a t g t c a t t t t a g c t a g a c a a c t t a t t 360
t t c t t t n t t t c t c c t t c t c t c t t 382

<210> 925

<211> 386
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 925

ngttcgagg tacttacccg ttgaagatcg aagaacgatg aataacgaat gaagaacggt 60
tgataacctt gcgagattcc tcacggaaaa cgttacggaa acgtttcgga agtgcctcg 120
cttagatntt cttcacggaa acaattnttc caagcaaatt cgaaggagag agaagtgcct 180
aaggggctgg acccctttct ttttcatttc ctccccattt tatagcaaaa tagggaggt 240
ggttgccgcc cagctcgccc aggcgagctc agctcgccca ggcgagcagg gttgcttcct 300
ccagaagcaa ccgccttctg gaggaatatt ccagagggcc caagtgggcc tgggtgctat 360
ttgcacccnn cattttacta agtaca 386

<210> 926
<211> 419
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 926

cgcctatnntt tgtttaatat gtntnnnttt tcaataaatg gaacttcatt ttatacataa 60
tttgttattag tataactatc aagtttcaac tattagaaat taaactagac attataactt 120
ttaaagcagt tactattata agaattaata tttttcata atatatacgat atccatgatt 180
agtttacagt atacaaaata tntatttcat taatataattt caattaaattt cttgataaaat 240
aaagacacan ttttaacatg atctatcggt tatatgaaag tgtcttcggg cagaatataa 300
ctctaacaaa atttctaaaa catagatata tacaatatca tataaaatataa aaaataataa 360
aattttaaac tacaatggca aggtntataa tgttataact tcgggttaaca aaaaaaaaaa 419

<210> 927
<211> 438
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 927

ctggtttaat gaaacatgca gnccatgtgc gccactggtg cgagtagaaac ctgaaagtgt 60

ggcgtagctg cttactgtga aggccgcgaa actgtgatgc cacaaacttc aagtcatcg 120
ctttcagggc aaaggcttcc acttctgaaa gggttggac agtcctggtt gaggtangaa 180
ggttgggtga agaatgagga tccaaagccc acgtgagaag ctccctcca cagaagtcac 240
cagccttgag gtactcagag ttgaagaagc cggttcttcc accgttagtt gtcatggta 300
atagcttgcc acgcattatg aagagcatct catcaaccgg atctccctcc cggacaatgt 360
agctttcttc tgtgttaagca ctggcttgag aaagcgcaca ttgatncaga agtgttcgtc 420
atttctcaac attggacn 438

<210> 928
<211> 491
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 928

tagtagcagt taccagaagt caatattgtg ccaatgacta tcatttc tatcttacct 60
annttgaaatt atggccttgg tttgtgttc tagattgaca ccagactcct aaatacacaa 120
tatcttcat tgcaagctta gcaactgtcc caaaacccaa gtttattcga aaccaagtgt 180
catgatttct atattaccaa ttntgctagt tgttaatgtt gaatcatagt tttgctctct 240
catctgccct ttgtctcatc tcatttacctt acaacttagt caattctatc attaccctt 300
ttcaatatgc agaatcagca acatgcaaac atatctaattc cagcaaatgc caccatcaat 360
agccaggcta tggccagaa ccaacaaaat gcctcatgtc ccatttctt catcttctaa 420
atntatttggaa gcttctgcag attaaaagaa gcatngttc ttcatttcac atgaatctac 480
tgggttagtt a 491

<210> 929
<211> 348
<212> DNA
<213> Glycine max

<400> 929

agaagaagtt catagagatt gattggattt tcagaaagat tgaattgatt gaaaatgcaa 60
aacaaaggct tgctttata gactcttcgt gtctggtaa gaagaccact tagaagagtt 120

ataacttta gaaaaactta aaaccaattt gaaaaagtca aaacctttt gaagagttac 180
atctttgat ttattcagaa acaaacactg gtaatcgatt accaaattag tgtaatcgat 240
tacacaaagc ttttgtgtga aaggatgtga ctttcacat ttgaatttga atttcaacgt 300
tcaaaggcac tgtaatcga ttacaaaac attataatca attacaac 348

<210> 930
<211> 408
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 930

ctgatcanat gactaanatt aatcgccaca acagatattt agcagtagga gggaggcctc 60
caacatacac ccgcctagca tatcatgtac cctgtaaatt caatacataa agacaggcca 120
taatcagggta aagtcatgct aaactaaaa taaaacttat ataagactgt acttagtctc 180
ggcatgcctc catactcgga aagactaaac taaaacgtt ttgtgttgag gaagtttagtg 240
tgtctctact ctgtatgtat aatgactctn ttcttcctaa taaaagagaa tatcttcttc 300
cagtagcaca atgatactat acaaacaaga gcgcaatata nagaacaca tggtaaaaga 360
agaccacaca ccttcatct acagcagcac atgatactgt aatatatg 408

<210> 931
<211> 455
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 931

agcttctcac atctgactct ctagttccaa cggtgttt tctttgatg cacttcctcc 60
gatcacccctg accaatggaa tctccttccc tcttaggtgc ttgttcgcc tatcttgc 120
cctcaaaggc aatgtccat atgtcaagtt cttttcact tgtacgtcat ccaatttgc 180
cacacgagat ggatcatgga tataactcact aagttgagac acatgaaaga caatgtgaag 240
gttagaaaga gacaggggta atgcaatttgc gtatgcccac agtaccgact ttttttagaa 300
tttggaaagg acagataaaa tgaggtatga gttattngaa tttcaatgct cgaccaactc 360
cagtccacaa agtgactctc aagaatacat gatcaactaac ctcgaactcc aagtcttcc 420

tccttcttgt cctgatagct ttctacctac tctga

455

<210> 932
<211> 328
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 932

tctacttcat aacccttga actacttcac attgatctat ctggccctc tagaacaatg 60
agtttgggtg gtaattacta tggcttagtt atagtagatg attactcaag gctcatatgg 120
actttttttt aaccaaaaat gaagctttg gtggctttta aaaacttgcc aagggtattc 180
ataatgaaca aggtctcaac attgttcac ttagaagtga tcatacgatgtt gaatntcaaa 240
atgagtcctt tgaaaaactnt tgtggagaaa atgaaattca ccataattn tttgccccaa 300
gaacacccca acagaatggt tttgtgga 328

<210> 933
<211> 445
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 933

ägcttggaga ttagtgcctca atggaggata agtaagagag aaggggggag cacgaaattg 60
aaggaataaa agagggagag aagttgaact ttgaagtgtg tctcataaga ctttcattca 120
tcaaagttac aacaagtgtt acacatgctt ctatattatag actaggtaac ttccttgaga 180
aaacttcctt gagaagcttc tttgagaaaa cttccttgac aagcttgagc ttagctacac 240
acactcctct aataactaag ctcacccct tgagaagctt cttgagaag attcctaaag 300
aagcttagagc ttagctacac acacccncta taatagctaa gctcacccca tgccaaaata 360
catganaata taaaaaaaaag ttccttattac aaagactact canaatatcc tggaaatacaa 420
gggtaaaacc ctatactact agaat 445

<210> 934
<211> 458
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 934

nttaactgaa tntgtacgt tccaaatgtt ttttanatgg tgtaatcgat tacaatata 60
tgtaatcga ttaccagtgt atctgaacgt taaaattcaa attcaattgt gaagagtcgt 120
atctttcat aaaatgctt gtgtaatcga ttacatggtt ttagtaatcg attactagt 180
acaagtttg aataaaaatc aagagatgta actcttctaa tggtttctc aagattctct 240
caagttata actcttccaa tggtttctt gaccagacat gaagagtcta taaaagcaag 300
accttgactt gcattgtaag aacttgatata aactttntac acannatttt gaacatcttc 360
ttgaacttct tcttcttctt ctttgccaa aagcttctta agttatttgg tttcaaacct 420
tgttttcac aaanacaaaa gtgtgtatta tctttta 458

<210> 935
<211> 401
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 935

agctnttctt tacaatcaat ctgtctgcta actaacaatt ctaaatgcaa gttcacattc 60
ttgttcttcc tttgtctaac atacatacta gctcaaactc atgaaaagaa acacaaactc 120
catcaaaaatc atgcactcaa tttaaagtact tttttttt cgtgaggaa aataacttgta 180
cttgggggca tgtcactcggtttggaaactc cttgtgact cgggcttatac accattgggg 240
gtgggggtgga gttgcctgtg cacaacagga tgaccttgac acttgctacc cagcttctt 300
gggtgtgagt gtcgtgtggg aatgctcang ctatccatg acgaatggta ctacattgca 360
ttttagagtt aaggtcaagt gcatgcatca tactaaggcat g 401

<210> 936
<211> 422
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 936

tcacacgtta taatggagaa cacatgatca gcgcgttggca atgacattca tggtaactccg 60
aacaatgggtg gagtatggag gattgccttg agggtccgca cttangcaat catgaaactc 120

agctccaaac tcgaaaagtgg aggacacatg aacagcccta agcaataaca ttcatgtggc 180
tccagaaaag gatgagaatg gaggattgcc ttgagggtcc tctcttangc aatcatgaaa 240
cacaactcca aactcaaaag cgaggacac atgaacagcc ctaagcaata acattcatgt 300
ggctccggan aaggacaaga atggaggaat gccttgaggg tcttctctta agcaatcatg 360
gaacacagct ccagactcga aaatggagga cacatgaaca gccctaaagc ataacattca 420
tg 422

<210> 937
<211> 506
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 937

ntttgctcat ctcgtccagn gatccttgag tcacctgcng catgcagctt gaagaaaaat 60
tagtattaat gtatgtaatg tataatntag tggnaatat taaggctata ttaatgtatg 120
tatangattt cattagaatt agaaaaaggg gtaattaacg tcataatagag tctaaaagtg 180
gagggcattt ttggtaatga ctatacaact agttaaaaa taggattta attaattaa 240
tttgtgacta attaaagtgt ctaattatta tgatgtaat aattaanata agttagagtt 300
gaacaccctg aanattataa ctcagactga cataaaactc tatgtnggc atctgtgtgt 360
gtatgaagtt aatttcagta gctataccgt ttaatcata gaantntcgt gctatgatat 420
atgtatgtga ctggtttagt aagcttgact gngaatagaa ctacctttgc tagattcatc 480
agtgcacatt tgactgtgat taagcn 506

<210> 938
<211> 397
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 938

ngtanggtta aagtctcacg atagtcacgt gctcatgcaa cagttgttag ccatggctat 60
acgagacatc ttgccaaaca aagtcaagtt agcgataact cacctgtgct ttttcttcca 120
tgctatatgt agcaaagtca ttgatcctgt caagtttgat gagttggaaa atgaggccgc 180

aattatacta tgccagttgg agatgtat ttccccctgcttcttgaca tcattgattca 240
cttgcattgtg catctggtca gagaaatcaa atgttgggt cctgtttatc tatgggttat 300
gtacccgatt gagcgataaa ttgcanaaga agccattgaa ttttttcag aataacttaga 360
gaatngctaa acctgtggcc ttctgagtct cgcatga 397

<210> 939
<211> 422
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 939

agcttgtaac ttatataata tatacatttt attgttaatta tattttaaaca catcagaatg 60
gtgcgcccatt aacccacagg tcccaggatc gaaacctgggt tctgataaaag agtggcttcc 120
gatctatcac atatatataat attttatgcg taaaacatat atcattacgc aatgacattt 180
gagtataata aaaaatagtt ctgcagggcc taacattca gtgcttataat taattttagtt 240
accatttaaa ttttattttt gagtcaactt ttaacgtat attcatattt tctctttgggt 300
aattntattt taatggctt aagtaaacat atttttatg gataataatg gcttccagtt 360
tcttagtgaa ccacatctga aaaattatac ttgaacaaga agatgtgttc actatgtcat 420
ag 422

<210> 940
<211> 427
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 940

ggtagagtcctntgtgtct cttagcttac acacatgttag nggaacatct atgnaaacct 60
gcaacgtgtta agcattgcgt tagctataat ttatgaagaa ccacttcttag ttctataatt 120
gtacaacata ttagcatatg ccaaactatg tgtatcattt ggatcaccaa aataagaata 180
ttacctcaat aaaatctcct tttggcatta gtgtctgca tgcatctcta ntcccttggt 240
atggtgatataaactatgt atgcaaataa caccagcatc tgcaaagagt ttagccacct 300
cacctganaa ttntatatcg tgatgtctaa ttatataaa aacaataaaaa tataatcgga 360

agatatacagg gaaagcattt agaaagcaac ataagaaaaa acagataaac tcaccaatcc 420

ttctaat 427

<210> 941

<211> 119

<212> DNA

<213> Glycine max

<400> 941

tgcttgagag acttctatgg atgttggatc tttgagcttc actatatgtc cttcaatgg 60

gattttcaat catggagttg catcggaga taaaggagaa gagggcgagag gaggtgtca 119

<210> 942

<211> 473

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 942

agcttgtgtc attaagcgat attccactnt tggcggttag cacgacctct tgtattgagc 60

gaatttacctc ctgggttgc aattgcactt agcgcacagg tctcgtaag aaagttgtcc 120

aaagatgtta tttgaaaaat ctcaatagta aaaaatgtag gcatgaatca agaaagttgc 180

agttcatgtt tgaaggtgat ccaacggta acgagtctgg gatcatggtt ttactgaaat 240

aggtaaaca aactccacat aaccttattt ttcacaccaa gcaaccgcac acaaataagt 300

cacacaacac ctcaactaat ccaacttaat caaagaatgc aagaattata ttaaacatct 360

attttcagtt atcaatattt taggctgtta caaaagacct tttcttggt atcaacacca 420

aagtattcaa gaatcttggaa gatcagaact gcatagggac acctganaaa aac 473

<210> 943

<211> 357

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 943

tagatactca agcccattca aatgaaagga attcataatt aaaaagttga gaagggttca 60

cngaagagaa tntcaaaatg actctgaatt gttttgtgaa caaaatggca ttaatcgtaa 120

ctttccgct ccaagaacac cacaacaaaa tggattgtg gggaggaaaa aaaagtccct 180
 tgaggaactt gttagtgttt agctctactg agctntaaaa gattggctaa gatcttgtt 240
 aaacataagc acttagacaa tgaatgaaag ctggagttgc tgcacatgtat gtccaacgct 300
 atgtcaagga ataagatncg gctgcacaat gcacaaggca agataaaaatg tcaaatg 357

<210> 944
 <211> 437
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 944

tatctctagc atgcattgta tgttggtctc gtccttgtc acgggaagcc ggaagggtcca 60
 tatcacccccc ttaattgtac acatggggca ctggcccccc aaatgcgcga agtagaagag 120
 ataattttcc gggctctcggt gtccgtaaaa tgcattcata tcattgcata cataaggcatc 180
 tcttcataac atcataatgg acatatcctg catttgcgtt ccatcatatt ccagcctcac 240
 attttgcattg agtcatggca tcattcatgca tatgcgttca acaaactttn tgatctgcaa 300
 aattgcatac catttgcgtt catgtttgct catccttgcg ttttcctcta caaaacaaaa 360
 acaaagaagg gggaaagcgtg aaacttcaca ctacattttt agtttcatgt gttaggagcc 420
 aaccatgttggatcat 437

<210> 945
 <211> 349
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 945

ctaacacctat catctctcat agtctntaga tntngagcc aatccaatcc ttgcgtccag 60
 actctcagcc acttatgata gccgccatg ctcccattac tgcttccctt aagctctatg 120
 tcctttcttc acgccgcattc ccatgccttg cgaactcctt ggagtaccct cgcgttgcgt 180
 tcactgaaac cccgtgcgtat gaaaggcgtg atgcttcgtt ctgatggcac tcctctcatg 240
 gggtagccaa gctgtcttat ggcgaggacg ggattataat taatacaacc ccttgcgttcca 300
 tcaaggaaac atttggacat ctttcgcattt aagatagaat cctgattct 349

<210> 946
<211> 156
<212> DNA
<213> Glycine max

<400> 946

cattctctct cattatcata ttagcattgt aggggggttc agagcattta tacttcttct 60
gtatctcgag gaatggtcta caaaccttga agcctagcgt tagttgtctt atacgactaa 120
attctgtata gaaaaacctt tgtcacagca tgtata 156

<210> 947
<211> 387
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 947

atctaagggg ttcaaaatgg aattgtgcat cgaaaactag ttctcaaaac ttggatacag 60
tagtgtagaa gtaacttata gaaaagagat agttatatag aagagtaaag aaaacactaa 120
ggttttatac cggttcacct caactcttgg gctatgtcca attgtcttcc aaaccttgaa 180
gggttccatt aatcaattct ttgattacaa tcaggtattc tctatgtcac ttctggctat 240
aatgagtact ttgttaccact catggcaeta cccttaatct cctcatgagt taagacttaa 300
gtattcttg tcactaagtc attccttagcc ttcacaaaca atatatgttt gatagaaaat 360
gattctaatac actcanagag tgttaca 387

<210> 948
<211> 400
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 948

tcacacatac acacaacana tcanagcana acaaacatcc aatcactgta taattacata 60
ctacaaaagc ctccataaca taacatcatc ctcaattctc aaaccctaaa caacaaagca 120
cccccgccac caaacacgca cacacaagca gctaaatatc attcaagaag taggatattg 180
ccataatcca caaacctaag aaaccacaaa actaaaccaa gcaccaacac cattccaaaa 240

acacacctcaca aaacattcac acacacacaa caaaccaaat aaaaaacaca cccaccacca 300
attcacatcc aaatcccaat caaacaccaa tcaataacac caaatcaac ttccaacaaa 360
tccaagccaa canacagcca tcaactacta canaccaact 400

<210> 949
<211> 384
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 949

cgccagcttc ttatccaagg ctcatcttgg tggtaagct tcttcttcca tggcttattc 60
cctagtggat gacgcctcct ctcaccttctt ctcccttgct ttccgctgca tctccatgg 120
ggaaaatcac cattaaagga catcatgaa gctcaaagat ccatcctcca tagaagcccc 180
acaagcaagc ttccatcaag tggtatcaga gcacaagagc ttcaagttagg tgctccttaa 240
acttccatta attntttgc tttaccttctt cttccattgt tggttcttca ttntttctcc 300
atgtatctcc tcacatgtct tggctaaat gttgttaaca tgattctta gagtttccac 360
caattaaact tgctatagaa gcta 384

<210> 950
<211> 341
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 950

tttctacttc ttccctgcca ttgtacaata aatgttagctt atagcatatt ccatagaaga 60
atctcatggc tacttatatac tggtaatgt gatattcgac aaaataaaagt ttcccattgt 120
gtactttaa gctaatagtt gaagacactc attgtgattt cggttgcgt ttatctcctt 180
tatgttaat tactcatttg acccctatac ttatagaaac tttctctttt agtccctata 240
cttaaaaaca tccctntta gtccctacac attccatttt tattcccttt cagtctctac 300
acatcatntt aatcccttgtt agccgctatg gtgaggacta a 341

<210> 951
<211> 464

<212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 951

 agctntngng atacttggtt catagcaata ttttgttgc cttgaaatga tatttagca 60
 gatatgtat gcagctctgg gatctttga ttccctttag tttcaatatt tgtagcatt 120
 tactactgtt ttgcataat tttgtttct gtgacccct tataatgtgg attaattcaa 180
 ttgagcaaat ctgttcgttn tatgctcagc ttctataagt ctttcattt acagatata 240
 tgtaatattt tatttagttt cttggttaa aactagttat tgttattgtt tctacccttg 300
 atttaaaact agttaaattt ataagatact ataattcaaa agggcagata gatccaaact 360
 tgatacacat ttcaatttga gagaatccna gtctgtgtt aatcatcgcc ttccatttca 420
 atgggattag gtgatggttt tactcatgtt atgactctat cata 464

 <210> 952
 <211> 309
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 952

 tgagcttaat gaattaattt attgattgaa cctggaggct attcagttgt atcttctgct 60
 accttatttt aagttttagg agagcatcat ccacagaaga tggttcaagg aaaatttgc 120
 ccaaatttgg gggaggtatt atcaacgtaa atntgttcca aatttgggga aggcaactcgg 180
 taacgattga aatggtcaaa gaaaatagta tatacacact ggctctttaa tctgtgtttaa 240
 aaaaaaaaaacca ataaaaaaact gtacgtataa ataaagttaa taagtgtgtt tgctataaat 300
 tcaggcatg 309

 <210> 953
 <211> 481
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 953

 ggatccttaa gcactggct gcagcttaag aaaaggccaa actctccctt ccaaaatttgc 60

attttaggct taaataggtg gcttgttcg tgcttgagcg cttagcgcaa ctctgaaccg 120
cttagcacga attagtgaat ttccggcttaa cgccgtgcctt tctcgctcag cggatggact 180
aaagcggtgc gcttaacgag atgaccctt gctcagtgaa catgcacagc tcatccttc 240
tctagattct tcctcgcgct cagcggatag ctcgctaagc cagtagattg gcttagcgag 300
aaggtaaaaa tcagcacctc acaaacttc ctaattaacc tgaaatttag agaaaaatgat 360
tattaaacac acaaaaatgga agtactaagt atttattacc tatcttaac aaaaagtaat 420
tacaacactt acaaataacc ataaattgga ggattntgtt acaattatgc cagtttatac 480
a 481

<210> 954
<211> 436
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 954

gggattccta gggcctaagt catatttggc actntattct agcttctaca aactgtccac 60
acactcaa an atgcagtcac atatgtacaa attntttcca caagctaaat tccatataaa 120
cacacgcaaa tgccatttag gcatgtcacc gaacacttga tgggtgcatt ttttagacatg 180
aaaaaaaaataa ggaacggngg gaatgtgaca tgcccattca tctcagagtt cacaataggc 240
ttgcggccat cccatacaac ccccccaattc aaacaaacaa gcatgaatcc aaacattcat 300
ttcctcatga aatntgaaaa tacaagcaaa caaagcacta aaatacagca atggcaagcc 360
aaagatcana ggagaatgac acttaattgt anggagtgga acaaaaatgca taaaggagaa 420
caaaaaactca acaatg 436

<210> 955
<211> 384
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 955

agcttgttctt ctctagcctc tnttngctt ttctacattt ccttagtgac aagtcatctg 60
cctagaagtt tcttgtcttg acctgggtgt tgggtcttcc tcatactgcca tgcacttcca 120

tgtttaattt cgagtgtctc ggtatattat gcgcctaaat tggacatccg agtaaaaagt 180
tatggccatt tgagttgcc tagaactttt gtgttcaatt ntgagcatct tgatatatta 240
ttggcctgaa tcggatatcc aagtcaaaag taatggccat ttgaatttc cttctgcttc 300
catatataat tntgagcgtc tcgatatgct atgcacccga atcggatatt cgagtgaaaa 360
gttatgacca ttgaatttc ttga 384

<210> 956
<211> 437
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 956

ntgaacaata tacttgtct tcatttaact gtcttgccc ttggcggcca cgctcaacan 60
agtactttcg acacctacta tacgttgatt tcaccaatgt tggtatggga atgttgcgac 120
aatccttan aaccttattt atacattctg agaggttcgt tgtcatatgg ccatatcgac 180
gtccttctct atcgtaagtc atcgccatt nttccttga gatgcgatca atccatgttg 240
ttatggctgg actcagttca cgaaattttt ctaaattttt atcacaaatg tgcttgcatt 300
gagtgtangc ctgcataaat aactatgaat aacactttaa gtttaatgaa gcaaacatac 360
gtgacatcaa tatgaatcta cccatttgtc acatttttt tgttacttat tgaatttcat 420
tgaagtcgct tgtatgt 437

<210> 957
<211> 525
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 957

atttttatga tgcgtctctg tacacagcca tcctctatag tcgcactgcn tcantgcaag 60
cttgaacaca cattgtgcgc gtgggtgata tccactacac aaggttgaa gtagaagaga 120
ccttctaccc tatcacacaa cgtggaggac atacgggggc ccgctacatg aatggccatc 180
atggcaatg ctgaaggat tctgcgcctc actatgcgag tgcacacaaat aatgcagatt 240
gtgcgtacgt gagcatgaac tactaccact atatagatgt gtggcacaca caagagcaca 300

tcttagaagc ttactccgca caatggtggc ctacttgaa tgaaggcagac atatctcctt 360
ttaatgacgc acggacactt atgcctgacc taactacgt tcacgcaaca tgtcagcgca 420
gatcaatacg gataangact gatatggact ggatcatacc atntgatcgc cgacngacac 480
gtaatacacg tggagcctaa ggggctacat gcatgctgcc tacga 525

<210> 958
<211> 523
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 958

tctaattaac ctacatatga gagaagatga ttattaaaca ctcataatga agataactaag 60
tatttattac ctatacttaa tagacaatac ttataagctt acaaaatagc catatattga 120
gagagtgtga tacaatttat acaagtgtta tacgtataag ttagatgttc tcaccgacta 180
acatcctatc aaggtcatca aaatttagacc agtttccatt cttgaatgac cctaacaaag 240
catgcatgta cgtgatcaag gtaaaggcat actagaatga atagctgata gcacagagaa 300
cacaccaaac atcattaaat agatagaatg atatttacat caagtaccta caaggaagat 360
ccaacagagg attntagctt tccatatcca ggaagccttc tttacaacan agagaagaat 420
aagatgacag agtgctgcta tacaagcggt gaggatgtct tcttcacctg taggatctca 480
caaccactca agaactcatac tcagactcat agaaaacggct tcg 523

<210> 959
<211> 304
<212> DNA
<213> Glycine max

<400> 959

atcggtgaga gtgtAACCTT aaactgtgag tgaacgacta gctgtgagta ataatcttg 60
catgaatctc tgaattttag aatgaaatgt ataactgaga acatgatgaa ggccatgatt 120
gtacatatac aagcttttt gaccaaacaa cttaccttga atgataattg catccttgc 180
tcccttttg agctgaatga tgTTgtaaaa aatttgaacc ctaaactaaa ataattatgt 240
cttgataacct tgTTtagatt ttaggagagc atatggtca aggcaaatct actctaaatt 300
tggg 304

<210> 960
<211> 344
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 960

tctggggatcatttgact tgctntccaa tctgacattc accacanatt ctgccttctt 60
ctattttcag aatgagaatg cctctaaca caccttgtc aatgatttc ttcatgcctc 120
ttaagtgcag atgtccaaat ctttgatgcc atattctgac ttcatcttct ttggaggata 180
gacatgtgga ggagtaactg gtttctttag gtgtccatag gtagcagttg tcctttgatc 240
tgctgcctt cattagaact tcactttct catttgtcac caagcattct gactttgtga 300
agtttacatt gaatccttca tcacacagct gactgatgct gatc 344

<210> 961
<211> 385
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 961

gtccatgagg aatcttcttg ggaaagacat cttaaattc ctgcaataag ggttgaacac 60
taggagaaac ataaatagtt tactgattag aattatcaact ctctctctt tgtgtatcac 120
tccatctctc aagtgtatca ctcttcctt ttctattcct ctgtgatgcc tcactattgg 180
ccctctcttg gtctctctt tctctcttc tgattcgac atcacacact tctctgaggg 240
ataaaagttt atgaataatt ttctggatcat ggtgctggag agaaatctt gttgagaacc 300
catcatgcac tgctttggag tcctcctcaa tcatggccct cacttagtgcc atctccatct 360
ncttatactt aacaatcaaa ctttc 385

<210> 962
<211> 247
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 962

ggtagaatgg ccagacatga tacatgtcag ggattggttt ggtcataagg taatatggat 60
 gtcccacatt attccatga cacatatgca aaaatgatga cttggatact ntatgc当地 120
 ctggcatgc atgcacctat gtggacactc atgtgtcaaa ttttatggt catgtatgc 180
 tagggctcag gattcattct ctctatTTTA atcaacccaa tgTTTCCAAA atatgttcc 240
 ttatcca 247

 <210> 963
 <211> 357
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 963

 agctttaaag tttaaaaaat gtaatgtAAC atttaatAGA gttcaaATTc ataactaaca 60
 tggcataatc aaactaaaat aaATAATAAC caagagtcaa taatcTTTC acgtgtcatg 120
 aggacgataa acctgagcat cattaacatg gtcctcagtt ccatctgtta tggtagtgga 180
 tggactanga gctagcatga tagactgaat atgcaagtgt tgttagtatat gctgcatttg 240
 ctcattntgt tgacaaattt gctcatntt tcttacatct gttgtcgcat ctcttgactn 300
 tggtggtcat gagtctccat tagtgcaac atctttcct caagtgtctt ctccTTA 357

 <210> 964
 <211> 406
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 964

 cgatgtacac taagcctcac atcttaggct aagcgatAT tgcagaaaAGA attttgtgt 60
 tgcagaaaAGC gctaaggcacc gctgctgcac taagccccAA atgcttattg gaagttacAA 120
 cttcaagttg ggcttagcgc ggngcttaggc taagcaccAG tggTTTAAAC tcaaATgtca 180
 tggTggcAcg ctaagcgcac catacgaatt tcagttttt AAAAGTAGAG gcagaggcAc 240
 ttgggttgct accttggcAc ccaaaccTCT gcactctcAc atctctgagc atatttctat 300
 ttctgctttg tgcttattga tcctctacat ctntcttcac actctgcatt acacaatcca 360
 agtaagtaac ttgatttctt ttcaCTTTA ttttcatgtt tcacag 406

<210> 965
 <211> 330
 <212> DNA
 <213> Glycine max

 <400> 965

 aaaaaaaaaatt aaaaattaaaa accagggtaa gggagcttac ttgggtggta cccttttca 60
 catttcctt ttcccaaatt tggaattact cttttatgc cgaagccggt taccggaaag 120
 tggctcgat cgccaagta ataattaaaa cggaatgatc cgagtgtcaa cacaggaaac 180
 ttattcctt ggcaaagctt tggcaacaa tcattgcattt tggtgacaga aaataataat 240
 tgtgaattga agtaaaagta tgatatatcc taattgaaaa gcagtaaacg tgagcaaata 300
 agtgtgaaaa cagtgtatcta aaagcattgg 330

 <210> 966
 <211> 421
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 966

 tgattntngt tctgattgca tgatgctagg atagttgata gttaanatag tggtaggaat 60
 tatattttca tacaatgtat gttgttctgg ttaggaattt atggcctaatt tggaaagca 120
 agcttcatga tgatgaacctt agcaattttt acgatgccaa aagaccaagt gattgattca 180
 agacttcaag atcaagcatc aagaatctaa tccaagattt aagattcaag agaagaaatc 240
 aagaagcaat aagtcaagac ttcatatagg ataagtatta aaagaatttt tcaaaaaacaa 300
 aatagcacag ttttggata caaaagaatt ntctcaaatt nttaagttt ccagagtgtat 360
 tactctctgg taatcgatca cctgttatca gtaatcgatt accagttgtc ataccctaatt 420
 t 421

 <210> 967
 <211> 488
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 967

atccttaagt caccgcggct gcagcttata aactgtttt aaaactaccc catattttag 60
tttttagaag taaaaaacaa aaacagctt aagatgtnt gctttcagt tttgttttt 120
ctttaaaatc tacaaaatac aagaataaac agagaattt aagtttcat taatgaaatt 180
tattnaaata tcttctctt atcatttctg aaactttgt aagcatttt ttggatatg 240
attattncaa ttggccatga tggtatgtt gtgccttgct aattatgtt atgggtgggt 300
taatggccta ttaatatgag aaatatagtt tgagtgggt acactcactc agctgaagct 360
tcttatatta tcacacaaaa gccaccagg acatatgtat atagagaggt gcagactaag 420
caccttatca aacggctaca gcagagaana agacatgaaa ggactgacta ttacaatnta 480
caagaata 488

<210> 968
<211> 355
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 968

ttgaattgct ttntggcctt ctggngtctc cctgcctcca attaagactc gatccgggtt 60
gaaaagatct tggattgcag ttccctcagc aaggaattca cggttgaaa ggatttggaa 120
cttgattccc ttccattgt gagtcaaat ctctctatg gcctcagcag tttcacagg 180
gacagtggat ttctccacca caatctgtc actctgnag acatcagcaa tcatgcgtgc 240
tgcactcttc cagtaagtca aatctgcaac ctgtcggtc tcaagaccac gagttttgn 300
cggcgtgtng acggagacga acactatgtc agcctcatag acatgtttct caaca 355

<210> 969
<211> 483
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 969

cggggattga tgcatttgca ctcgagatcc tctgagtcac ctgaagctgc acgctctctc 60
aactactgga cacatggttc acagatgaaa gcaagaagaa tgactacaag atgattttatg 120
ttgtaaaaaaaaa tgtaaggatc accaagtatt tgaacttgta atggttctc cggttaagctt 180

caacttccc aatctactcg aagcacaaga ttgtcaaagt tggtgaaat gaaaggagcc 240
tttcatccac agctagtcaa gggtttctac agctatgcac ttgttgaccc ttgaagtaac 300
ctctcttcta aataaatgga gtaaaatagt ctgtatctat tgatggaaga agtactggct 360
ggatatgggg gagtcacaat tctataaacg atgatggac aacagatgag actatganga 420
tgttctgacc tttagaactg aaaatctaa agtgtggctg ttgagagcaa atgtgttatt 480
ctt 483

<210> 970
<211> 376
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 970

ntgtcctcag atcactcttg ttggactcag cccaatcgag atactcctct taggtttaga 60
ctaacttaaa ctgagttca tctgcagatc cctcttgtac caatgccaaa tagtactcta 120
acaggagatt ctgggtgga agtagtggat tataccatca agacaagaga gcagataaca 180
aagttaatta ctccaccaca acttgctcg cgcccaggaa agaatgaaac attatgctga 240
cctcaaacga gtggacaagg agttcaaatg tggagaccta gtctattnaa taattcactc 300
atacaagcag cttaactttgg canactatgc ttccacacaca tagcagccac tagcggctca 360
tgtaaagaaaa catcat 376

<210> 971
<211> 338
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 971

ngttatatac ggaagacaag aaaatggat atgaacgaa tgatgatgaa agcttagaat 60
ctagacatga attgaaagtc tcagattcga aaacttaccc gttgaataat gaagaacgaa 120
tgaagaatga atgaagaacg acggaaaacc atcatggatt tgctcacgat aacgtctcg 180
aagcattaca gaagcacctc ggcttggatt ttcttcacgg aaacaattnn ttttcaccag 240
aacagctgaa atgcatagcc agggatccg ggatccttgg aacaaccccc ttttctctc 300

tttataagaa aaggcgagag gatgttgctg cccagctc

338

<210> 972
<211> 393
<212> DNA
<213> Glycine max

<400> 972

aagatctaac agattcaacc taatgttcat gagaatgatt gttgcagata atattaagat 60
aactctccct aagactgata atgctaaaaa gtttatgggg ttagtgggag agcgctctca 120
aatagcttat aagtctcttg ctggcacatt agtgagtaca ttgaccatca tgaagtttga 180
tggttcacgt actatgcatt tacatgtcac ttagatgaca aacattgcaa taagatttaa 240
gacctcggga atgggttgta ataagaacctt ccttggtcag tttattttga actcattacc 300
atttgtgtat gacctgttcc aaatgagcta taataccatg aaagataaac ggaatatgca 360
tgaatttgcattt agtatgttag ttcaagaaga aac 393

<210> 973
<211> 277
<212> DNA
<213> Glycine max

<400> 973

aggattgatg gcgacccagt attgagagat acgaggatat gggctgcgtg ggagtacgtg 60
agctcagttg gcgggtggca acagggatg gtgggtttat gcgcgcattt tggatgagga 120
aaaccttggc tgccacatcg accgaccgccc acctagtacc acatgtgtat ggtaccccat 180
aatcctacaa gcttggatg aggaagtgtg gaagggtgaa acttccttgc ttttagttgtg 240
accacagagt ggtaccttggaa gatatgtcgcc gggggc 277

<210> 974
<211> 457
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 974

agcttctaga gagttctaca ctactctaat attctccttt ctacactntn ttaaaaaagt 60
ataaaaatttt ttagaacctc tccaaatcaag aaggaatccc aacacattcc ttaatgcggaa 120

tatcttcatt tcaagaagaa ttaagtgatg aaaatataag gttggcttga tgataaaggg 180
tgaaggaaga gagggatagg aatcttccgg ttaacaaaaa ctaataaatt aacaactaac 240
atttaccgat aaaaaaaaaa aagatagaat taagtgatta gtttatctc ctttcacat 300
tatagttct tcaatgttc ttccacaattt ctaaaacatt gtctgttagac tgctccaagc 360
ccagttcctt ctcaaacagc cccagagaaa tgacattgat gtgtctaata taagttctct 420
cttagtctta aaatttcaaa tgtagtgaaa actatac 457

<210> 975
<211> 577
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 975

attgaccat gtgaangcga tctattangc gacactacag aagactcagc tcagtcac 60
tagtcggcag acttatctga atctttgaa gatgagagaa cacgccaatt aanaagagaa 120
agggagaaac ttcaagaaca tggtctcac tttcacgtc caagagaaga atccccacaa 180
atatcacaaa caactcttcc acggattca ccaatccaag cactcacgtt caaagcaa 240
aacaagagca acagaatcaa acccatcatt gtctttgtt tttccatggg tgagattgaa 300
aaaaaaaaacaa caaanaatgg tatcttcac aagcactatg tctttggatt ctatgagtag 360
ttttcttata accataatgc tcctaaagga tgtttatggc tcatagtgc tggatnttt 420
caccgacatg atatcttata ttatgttaa gactatctga actntatgtt aagatcact 480
ggattctcaa tttaatgga tggttnagc taataaaaga atcttatatt gtacgcctac 540
aaataaaaaga acaaatcgtn taactttaca ttcttn 577

<210> 976
<211> 460
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 976

agctnttagc attntaccat attatataat ggctagctt gttatattc tcctaactac 60
atggcatgcc aagatatcac taatatgttt aatgtccacg gttgtttctg ttaagttgg 120

gctacgtgag ttgactctga taccattatac attatcatga tgtactgttc atggacccaa 180
ctatatatta tattattaat tactaaaata agagtttagtt ttaaaaacact acaactttc 240
tactataaat gtattggta aatcattcat tgaaatgata atgctactta catatattaa 300
ttatttaaaa atgtatatta taactntacc acttctctaa aaaaacaaaa aanaaaaactt 360
ttccattta cataaatatg taatatggtg ttaattgata tgtatcaagc atttattttc 420
actaacccaa gctgcagga attctatgta ctttgtcatt 460

<210> 977
<211> 307
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 977

tgcataaaatg agaggaaact ttagtaagtt ntatttataa atatataattt cagatgatca 60
tcttttaatt tcgaaaaatt atattcaatt ttattctgca ataaatgttt agttattttc 120
tggtaattt gaatttaatg ctgattttc aattaatcat ttctaaacta atatcatggc 180
ttgtggacat tggaatctat tacattagtt tccccacgct aaaaaaaaaa tagtgatgg 240
caacaattgc ggcgttattc catgtgcattt gagatcaata caccgacaaa atagtgaatg 300
cacccat 307

<210> 978
<211> 260
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 978

cttgatagc ctttntgag ctttgttcc cttcccttgt tttgaagctc actacaagcc 60
ttaagtgaaa aaccatgata tcaccatatac cttaaggaat tttggagctt tggaaattgtt 120
ttggaaataa gtgtgggggg ttttgggttc attggacaac ttgtttgtt ggctatgctt 180
catgtgtat tctggccat acttgatgta cattgtatat tggtaaatg ttggacatgc 240
tgaatgaaat gttgttctc 260

<210> 979
 <211> 401
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 979

tgaagggaag agagagatcg atcacgagca catagcatgg tcttaanaga agagttagca 60
 gcttgcttaa ggtccaaaag gaacttgact cagcgttgt gcgagacaga gatcaacatg 120
 ttggctatca tcaccaagta tcaataagaa ctaagtctag ccacagccc cggcatagg 180
 gtggcaaacg agtatgccc agtgtacgca gaanaggagg cttaggaaag ggtgatcgac 240
 tcgttacacc aagaggcaac catgtggatg gaccaattt ctcttacctt anacgggagt 300
 caagaacttc cccgattgct agccaaggcc aaagcaatgg tggacaccta ctccgcccnc 360
 gagggagatc acagacttct cgactattgt cagcatatga t 401

<210> 980
 <211> 363
 <212> DNA
 <213> Glycine max

 <400> 980

gcagctaact accatgcact acggatatac tctaaggaac gcaaaaagta acaacaaaga 60
 ccacattaag aactacatat gcagcgacct caacgcaaaa gtattacctt cactcctcgc 120
 aaccaaaggg aacaacacac caaaggaaaa atggaaaaca aaggaaactt atccaaaaac 180
 aaagcatgaa agtcagcaaa caacaaaaaaaa ggacgcgaaa agatagaaca agaaggaaaa 240
 atttagtaaa gaatcataga acgtgaagta aacaacaaaa caataacaaa ccgggtagaa 300
 aaacagctac agccgcgcca aacacagaca tgcacaacag acgtaaaaaaaaa cttacacctt 360
 gaa 363

<210> 981
 <211> 366
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 981

cgctcggtga gcttctatgg aggctggatc tttgagcttc aatgaagtcc tttaatgggt 60

attnccact atggagatgc agcgaaagac aaaggagaag aggtgagagg aggcgccatc 120
cactgtgaa taagccatga aagaaagagt ttcaccatca agatgagcct tggataagaa 180
gcttggagg atgcttaat ggaggaaaag acagagggag agaaagagat agggggagca 240
agaaaatcgaa ggaataaaaag agggagaata gtgaaacttt gaagtatatc tcacaagact 300
ctcattcatc anagttacaa caagtgttac gcatgcttct atntatagac taagtagctt 360
ccttga 366

<210> 982
<211> 514
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 982

tgcnttattg ctctatctac tcccngnac ctgagtcac ctggccatg cagctntcng 60
gtgcggtcta ggacacaatg tcaattcata cgatatgcga ggatgactcc ccgagcaagt 120
tggatttggt atgaccatgc ctcctggtt tctgactang aaattggcga gtggaggagc 180
gcccacacat ttacgcgaca agcataatgt aacccttgc ggctntaaa ctctacggng 240
ggccctangc nttagagatt cctttgtta tggcattatg tctttgttc ttgaatttat 300
aaatataaaag atcttcttc atctgttccgc acctctac ccattctcat tcatttgcatt 360
gtntatttct ntacgcttaa nacactagat ccaacaacga gtcctcnaa ggtactaata 420
cctgngaccc tgncatcgat tcatgcaaga agcgggcaca cagagagtga gaggacgatg 480
atgtgtactt tccccacagtn gagaaatagt actn 514

<210> 983
<211> 369
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 983

gcatttggac acctattatg tatctcctat gctgtaccta catacgatc agcagggcca 60
ccatctcaat attacgaga tcatattcat acaccattgg ggcatttcac caagcacttg 120
gtgagcgcatttggacat gaattgcaag agaatggag caatgtggca tgccccattg 180

cttcaaaata caacctatgc ctaagaccaa ttcattcaga ttctcaattc aagataaca 240
gcccctaaggc taaccataac tgcctcacaa atataatgca tggttcaca atttagggca 300
ccaaaagatg aagaaaaacac atcantggga agcatataca tcaaagatcg aataacttact 360
tgttggagt 369

<210> 984
<211> 419
<212> DNA
<213> Glycine max

<400> 984

agtttcgtg atttagttt caccgacgaa atgatcgaag tgggtctaaa aagaggcaaa 60
tctgatcatc atgctttgtat aaatgaaaa aaaattgggg caagtgaaga gggtgagaat 120
gaaggagaaa cccatgttgt gactgccatt cctatacagc caagttccc accaacccaa 180
caatgtcatt actcagccaa taacaaactt tctcttacc caccacccag ttatccacaa 240
aggccatccc taaatcaacc acaaagctg tctatcacac ttctaatgac gaacaccacc 300
tttagcacga accaaaacac caaccaaaaa ggaatttgc agaaaaagc ctgttaggatt 360
caccctaaat tccggtgtca tatgctaaac ttactctcaa atctactcaa taattcaat 419

<210> 985
<211> 423
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 985

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ntatttcgtat ntgatttca ttatgtttct ttaattntgt ttcttacgta aacaaagcaa 120
ctgccaaatg catttccctg gaacaaattt atttgaatat gtcttgggt ttgttaatc 180
aattnttaac nttagttaga cttacacaaa tatgttatgt tatccaatgg gcataatgaa 240
cggtctaaa gattagacaa cgtatattct tcattaagaa aaaggaaaag gcgattataa 300
ttntgactta agaagttgtt ntgattctgt ttactgattc anaagttggc tgattgttn 360
ttagttntct actggatttt attatccttg aggacttgc gtgtcctcac atagtgtatt 420

<210> 986
<211> 311
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 986

tctgttgttc aatttcgagc gtctggatat attatgttcc atattcanac atccgagtga 60
aaagttatga ccattagaat ttctcgagag cttccgttgc tcaattcaa gagtcttagat 120
gagttatgta cgcgaaatcga acatctgtgt gaaaagttat gaccattcaa atatctttag 180
tgcttccgtt gtgcaatttc gagcatcttgc atatattatg tcccacattt ggacattcgt 240
gtgaaaaggat atgaccattc gaatttctcg agagcttcca ttgtttaatt tcgagagtct 300
agatgagttat 311

<210> 987
<211> 575
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 987

attgcacgca tctattangc gcactatacg aatactcaag ctcttccgga gcccatgaat 60
cngcgtttcg ttcatgtgtc tccacccccc gagttggagc tatgcgttagg gattgcttag 120
tgcaattctc cattctcaac cttntcgga gccccatgaa ttgcgtttc gttcatgtgt 180
cctccaccct cgagttcgga gctatgcgtt gtgattgctt agtgcatttcc tccattctca 240
aacttttttgc gagccccatg aattatgttt tcgttcatgt gtccctccacc ttcgagttt 300
gagctatgcg tagtgatttgc tttagngcaat tctccatttca aaccccttta cggagcccat 360
gaattgcgtt ttgggttcatgt gtgtccctcac cttcgatgtt ggagccatgc gtagagattt 420
cttagtgcaa ttcttcatttca accccctttt ttggagccca atgaatttgcg ttnntcgta 480
tgcgttctcc accctctcgag ttggagctat gcgtatgtat tgcttagtgc aattctccat 540
tctcacacccctt tttcagagcc catggattat gtttgc 575

<210> 988

<211> 415
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 988

gtatatgaca tgtcccaactt gtactttntt ttttatcta atttgcatcc cacaaaatta 60
gaatctctgg atcttgattc atccactgat ttaccttct catthaagtc aaggtaggtt 120
gatgtagcca taagaatgga cgcttcttg cattnntcca taccaaattt ttttattagt 180
tctatgcagt atttattttg accgaggaag gttccatgtt tcattttctt gacttggagt 240
cctagaaaaga aatthaattc tcccatcata gatgtctcaa attcttttt gcataacaaca 300
tgaaaatccc ttgcataagg tttcatttagt atagccaaat ataatatcat caacatata 360
ntgaacaatt aacaaatcat tgtttacttt cataatntaa caaagtttgt caact 415

<210> 989
<211> 260
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 989

cgtatgataaa gactccccaa gctatntatc ttctctctca gagaggctnt gtctcactct 60
aagaagtgga tcactcttat ctggatgga taggaatgaa agtcctaca ctatattata 120
ctactccatc tncacaataa atggtgaga ttacttgct cataatgtga agattaattc 180
tctataatgc ttcacacatt ctaagagttt ctacactctt ccattttctt tcataagggtt 240
ccagaaagtt ttacacatct 260

<210> 990
<211> 181
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 990

tactttgtca ggtacatggt cacttgcaat ttcattcctt agtcctgtt cccttgc当地 60
gatcttccaa gatactatta taacttctt aggaatctct tctgaaggctt atccttcaag 120
ggtagcagcc ttctttttg cttttcttc tgctntcttc ttcatgggtt tttatgctt 180

<210> 991
<211> 585
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 991

agggtacaca gcacaaaagg acngctgcgt aangctacct atcattactc aagctcgat 60
gaggaagtgt agaacggta aacttcctgc tnttattctt tgaccacaga gtggcacctg 120
gagatatgtc gcgggtata gtcagtcagt agaacctgtg atgtaccta gcaggcgagc 180
tcctggcagt caacagataa aaggaacaaa gatcacaaag caaggaggct tgtgtggttg 240
ctggccagtt gtgaaacttg attgatatat gggatgtggc ctctggtaat cgattaccaa 300
gggtggtaa tcgattacaa ggcttanaaa gtgaagacag gaagctaaga tggcctctgg 360
taatcgatta ccaaggggtg taatcgatta tcangcttga aaatgggatt aggaagctaa 420
gagggcttct ggtaatcgat taccaagggg tggtaatcgat taccangctt anaaatgaan 480
gcagcatgtg gtggaggcct ctggtaatcg attaccaggc tggtaatcg attacacagg 540
ggaacatgcc actggtaatc gttaccaggt atgtgtatc gatan 585

<210> 992
<211> 320
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 992

atcctctgag tcacctgcng ctgcagctta catggagcta cgtcattgac ggtttctcat 60
attccttgct acgttaataa gcataacctct gtatgtctcc atcttgatgt agccttnat 120
tgattcctca aacttgcaga ctccccata ttttagtcctg caacttgctt tcaaactgca 180
acatcaatga tcatctctac atcattaaca gtagagttata aattgccttc catatcaaca 240
cgggcacatg tggtagaaaac ctttaccaac tcaaggtaga agattnntat cttctacacc 300
aatcttaaca aaccatatgc 320

<210> 993
<211> 126
<212> DNA
<213> Glycine max

<400> 993

tatatacatcc agttccagtc atggtatata gtataaaaat ttcaaacatca acacaacatg 60
caatgaagcc tagcttccaa agacaacaag gttagggttc aacaagtgga aagacccccc 120
ccccct 126

<210> 994
<211> 248
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 994

attcccttag caatccccca aattaagaac ttatcataac ttgaaaccct tatactctct 60
tagaaccccta aaacaagatc aaggttatca aaattaggct caggggtta ttcaaacaaa 120
tcattattac tttggctca atagggtgc aagggataaa ttcatcacag gttggctntt 180
tggctgagtg gctaanataa aaagaaacna tggcttgatc atatccacct tatgcaaata 240
atcaaata 248

<210> 995
<211> 305
<212> DNA
<213> Glycine max

<400> 995

tatagatact cagcctcatt ggagctttag gcctaggatc ttcttatcaa tggattctct 60
tgcttcttgg aagatgaatg gaagcgaaat ggagaaagg agagagagag gagacgccac 120
ttcaaagaga agatgagtct agaagaagct caccaccata ggaggccatg gataagagct 180
tggaggaaga aggagatgaa tgaagggaga gggagagaag agcacgaaca ttttgctct 240
acatgagctt tgagatctga agttaatat tcaaatgatc aaagttgaaa aaaatgcaca 300
cacat 305

<210> 996

<211> 362
<212> DNA
<213> Glycine max

<400> 996

gcttgttggc cgcgattgac aaagggtgca tatatacgac gttagtctct gcatgctatc 60
atgcgttgac tggtagcgat agcaaaaagaa tgtttatact aataaccact tgggtatttc 120
tgccggcccc ctaacttcac gacttagtac cgacagagtt tgtaagcgtg gaagacgacg 180
taaatctccg catgtgaacg agcttggc ccgcgattga caaagggtgc agaagacgac 240
atttgtttt tcattgttac atgcattgag tcttagagat agcaaaaagaa tgtttatagg 300
gataaccact tgggtatttc cgccgacccc caacatcacy agtttgtatt ggagaggttt 360
tt 362

<210> 997
<211> 416
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 997

nttaggacct tatgtttagt tnnttgctt ctaaatcttt tgacttgtaa acaaaaaagt 60
tctttgtat atttcaatgc ttaagtgaag tggtagatt atgatgtta caattacttc 120
aacaaggcct tcgataataa attgttcttt ctgttgcatt aagcataatc atgcatttc 180
ctgcattcat agttccgca tcaagtctca cactgtgttc accactcaa aaggataatc 240
agccgcccgc cgaagaaagt ggccgcacga attctccgca naccaactcg catttcaga 300
aatggatcta attgagcaag aaaatcagag tctcaaggag gaggttgcca ctntacgaga 360
aggaatggat aggttgacga ccatgatgaa tgcactcctg tccgcccaga attctc 416

<210> 998
<211> 435
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 998

agcttgaggc acttgccttt taaccttagt tctccaaagt ggccttaccc aagtatctt 60

ttttccattg atgtgccatc attttcttct attttctaaa ccctttgc accatttaa 120
ttattgattg gtcttaatg tcaattaatt aggtagttt attatttggg cccattcagc 180
caatttgatg ttttaatct aatttcagga attaatgaag aattgggctt gaatctaga 240
ttgggcttga atctagaatt gggcttggac ttgaagaggg caaactaatt tattctataa 300
aatttagatct tatcttatct agatattatt tagatttgat ctcatctaga tatcatttca 360
attagatctt atcttatctt atcttatcta gatntgattt gatntactt atgggcttgg 420
attnaaaaaca tattt 435

<210> 999
<211> 243
<212> DNA
<213> Glycine max

<400> 999

tgacactact tatcttaccc tacttctacc accaaaatta agtataacct atagaatttt 60
actcctgaat taattaatta aacgaatgtg tatagaactt tctatttct tttcataagt 120
aaacattcct cgcttagacg ctgcgtatc ttcatagcgc tccaatcata ttaatagtt 180
ttccacccca ttctgtgata tacaacctga aaagctctga atatgctga tacggaatta 240
gtt 243

<210> 1000
<211> 257
<212> DNA
<213> Glycine max

<400> 1000

taatttatct caccctcatt tgtcacaaga tagtgacatg gagttgatgg tcaatattt 60
tcacaacaaa gtatgtttat ctcaacctaa tttgttgag cactccattt ctatataatta 120
caattattca tgttcggcat tagcatgtac gtccctgcaa ctattgttcc acccatagca 180
aggataataac taaccataac atgagccaa caaaggaaga atgctgatat agatgatgca 240
gtataaagaa aactgaa 257

<210> 1001
<211> 522
<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 1001

ntanatatnt cagatgaagt cattcaatta ttgttacttc atatgttagca gctgggagga 60
gatttatatg tttgtcaaat gttgttgaca atgtagaaac attgttaggc atgacaaccg 120
gatcggttgc tgcttatcct atctcgagtcc ccgattcctc atttccttct ctacccctgt 180
ccccgaaatt caatggaggt gcataattat gtccatccca gtctccagtg gggttgagtt 240
tttcccgtcc cgtcctgccc ccgacatatt tataaaattn tattaaaaaa tctaattntt 300
cataaaatga agaatataga tttaaataaa aatcacaata ttgtacatga caacatanaa 360
tccaattcaa cattagcata naattcaata taataatttc atggtttaat gtgttatata 420
tatatatata tatataatga tatnnttgta caataattat tagcgtggag gaattggagg 480
cgggtattaa taatctcatc cctgaccccg aacctgattt tg 522

<210> 1002

<211> 267

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 1002

ctcgctgcan aattcatttc ttgggttgtt ctcttggtt gtgctaaagg tgggtttcg 60
cattggaagt gcggcacaca gactttgtgg ctgacttaag gattgcctt gtggataact 120
ggcggtggg taaggagaac gtttgttatt ggctgaatca tgacatcggtt cggttgggtgg 180
gaaacttggc tgtctatgaa tggcagacac agcatggct tcacatccat cctcaccctc 240
ttcatttgcc ccagtttct cattcg 267

<210> 1003

<211> 360

<212> DNA

<213> Glycine max

<400> 1003

tattccgagg acagttcatt atcatgcaca gcctgcaaga gttggctcac aatacgccaa 60
tcatatctat ggagcatttt ctagagccag tagcctcgcc tgaagctcaa cttccattgg 120

tgaaacccaaa cgagggttgc ccgcctgagc tcacacctga gtaggtcaat tcagagccag 180
ctaaccaca atctccagtg gcgaatccac cttcttcgct tgagcttcaa gcaagtcccc 240
catctcctcc tctgaatgtc atttctgacg catcattaga tgaagcattt gtccttctg 300
atttaccagc tgcagataca gctgaccacc ttgtttcccc aatcgagga catgctgatc 360

<210> 1004
<211> 237
<212> DNA
<213> Glycine max

<400> 1004

agcttgcttg agaagcttct atgtaagctg gatctttgag cttcaataaa ttccttcaat 60
tgtgatttgc agccatggag ttgcagtggaa agataaagga aaagagatga gaggagacgc 120
catccactag agaataagac atggaaagag aagcttcacc accaagagag tgtcttggat 180
aagaagctta gagaggaagc ttcaatagag gaagagaatg agagagaggg aggggggg 237

<210> 1005
<211> 243
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1005

cttganatcg aanactaact gggtgaagat ngacgaacaa tgaatatcga tataagaatg 60
gtgaagaaca ctatcataat tcatcacgaa aacgtcacga aagcatctcg gcttggatta 120
ttttcttctt tcttcttctc ctcactaatt gtaagtgaat tttgagtgcc aaagggtttg 180
aaccctttt cctcagcccc ccatgccatt ttattgaaaa aattgagggg gggggggggt 240
ctc 243

<210> 1006
<211> 315
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1006

gcttcctcca caagcaacag cttctggag gaatcttctg gaaggccaa gtgggcctgg 60

| | |
|---|-----|
| ttgctatttgcaccccccattttactaaatacaccccttctattttttggtgattctt | 120 |
| tttccgtaatgttacgaaaccttacgaatttcgtaacgactttatttttcttccgcaa | 180 |
| ggttacgaacccttacgaattatgtatttactctntttgccttcgaagaagttacgga | 240 |
| aacttaccgatttgcgc当地acaccccttttcgacttccgtcacattatggatattcacg | 300 |
| gatcgcgcaagcctg | 315 |

| | |
|-------|-------------|
| <210> | 1007 |
| <211> | 446 |
| <212> | DNA |
| <213> | Glycine max |

<223> unsure at all n locations
<400> 1007

| | |
|---|-----|
| tattatggtc taagctggtt acacgaacat ntattcccat catatataat aaataaaaac | 60 |
| aaaaatttagg ctcccaccaaa aacaatagtt tgaaaatcaa cttcactaat gttggagaag | 120 |
| tcagagaaaag ttgcttagaca aaaaatatct ttgtcatata gctggaatga tcatccgtga | 180 |
| gagaaaattc atggcaagca tagcaagggtt cattcttctt gaaaccacaa ttggaagctt | 240 |
| caaactaata atcttgagaa ccatattatt taagcttcaa attaatgcat cttaacttta | 300 |
| gtcacgacta gtacactaga gacacgataa agtcaccaga naacacgcaa aagtacttaa | 360 |
| taaanactta taagttaatg gtttagattaa taacatttag tttaacatgg ttttaggttg | 420 |
| tcaaaatttgt ccgtcaaaaa agtttc | 446 |

| | |
|-------|-------------|
| <210> | 1008 |
| <211> | 497 |
| <212> | DNA |
| <213> | Glycine max |
| <400> | 1008 |

<400> 1008

| | |
|---|-----|
| cgcatgatac atctgacccg cgatctctga gtcaacttgc agctgcgcag cctggtgccc | 60 |
| catgaaggat ggcttgcggt tatattgcat gaaaaagccc ttcgattata tggatatatg | 120 |
| tgaatgggta gcataaaaatg gcttgcgaaa tggtgaataa aatggcttgg caaatatgaa | 180 |
| tatatatattgc ctggaaatgg cttggattat atgaatatat attgtatgaa gtggcttacc | 240 |
| aagggggtgga tggatagccg aaaagtggtt ttcaaaaatat gtggatttgt gaagagggag | 300 |
| caaaagaagc cttccaaaaa aatgtgtgat atatatagga tgtaacgtga aagggttgc | 360 |

aaaaatatatga catggatgtg tgtcgaaagt gcttcacaa attttatgtg tgcaatgata 420
tgtgtataaa atacatggcc caaatgtat ttataagtgc tgtgacactc gccccatgag 480
tgtgttgct cttgttg 497

<210> 1009
<211> 441
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1009

tattaagaag cttcctccag aagcttcctc atggcttctn tgagaagctn tctcaacaag 60
cttctttgag aagcttagatc cttatctatc cacaccctc tattaactaa attaacttcc 120
ttaaaaataaa ttacggatga aaataatgca acaaataatc aaacatcaaa cataattact 180
aataatataat atatcagggt gttacacatg gtatacttga gaccgtatag taagcataaa 240
atttagtata ccaagaacaa tgcctttta ttgactacaa ccaaagctat aagggtcgcc 300
aatgataggc actaagttgt aagatcaata tttctataca tggtaattt caagagttgt 360
agttccttcc taaaactanga acaaaanana aaggataaaa aacatgccac ccctctaaaa 420
tatcacacacaa ctntntttaa a 441

<210> 1010
<211> 444
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1010

agcttcttagc caaatattag tctctcaaag agtcgtctt tcatggagga ttatctgttt 60
ctagagtctc atcaattcga ggtgtcctt gtttatgga gactcatttc tatttaacta 120
tcttggagtg ccaatcttc ctggaaaatc gagaaaata tatctccaaa gtgtgtttgg 180
atgagaaaat tttaaattct gagaaatttt aaattctaag aatttcaaattt acttcaattt 240
aaattctttt attttaaaaa ttgtgtttgg ataaaaaaaaa ataaaaattt tgagggtgaa 300
agaaaaatgaa tgcaaaggga agagaagata tgattgggtgt gttttaaag agaagaatatt 360
tgacacggca tggagagtca cacganaact gggacacgac gacatacacc accatacccg 420

accacaacat tcagtcaatg acac

444

<210> 1011
<211> 470
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1011

tactcaagct tcaacaacga ggttcagtga atgattacct cactgagtct gaacgtttag 60
cgaatcgat tgggtctg tctccacgg tgctcctaag ctgtttatt tcgggcctt 120
acccagacct ccgtcgtgaa gtccgggctc tgcaacctat gtccatatca caagtcgtgg 180
cgcttgccaa gttgtaggaa gaaaagattc aggaccgcca tcgccatttc cgcacatcct 240
atacccttc tggccgcca ctgtcaccgc caccacccac cgcggttctt tccatcggtc 300
tcaccccggt acgccctta gttaaagcgcc tttcagcaga agaacttgggt gtctgtcg 360
acaagggtt atgttatcat tgtgacgaga agtggattct cgacaacggtt gccgtcctcg 420
cctccactta cttattgcan acgatgatga tggatgactgc acaaatccat 470

<210> 1012
<211> 417
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1012

ttggcttatt cccttagtgga tggtgccctcc cctctcctct tctccttgc ctgcgcgtgc 60
atctccatgg tgtaaaatca ccattgaagg acctcattga agctcaaaga tccagcctcc 120
ataagaagctc cacaaccaag cttccatcag gacgaagttt ggattgattc aatctaacta 180
gggattgagg tttagtaatt taagctatag catagaacac aaaagcatga tngatttagag 240
aaacatctt atatacatca gttgggttgt tagaaagact caacatctt acctactggc 300
tgcaatctta cttactttgc atttactgg ttttagccta gatntagtn aattctattc 360
taaatcatcc attatcaatg gttctctcac aatgacttat tctgaattaa ccctatc 417

<210> 1013
<211> 391

<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1013

tctcaaggga agtttcttaa gaaagcttct caaggaagct acctagtcta taaaatagaag 60
catgtgtaac acttggttata actntgatga atgagagtct tgtgagacac aactcanagt 120
tcaacttctc tccctttgtc ttccattcaat ttctgtctcc cccctctcta tttctctccc 180
tctttttttt cctccattga agcatcctct ccaagcttct tatccaaggc tcattttgg 240
ggtaagctc ctcttccat ggcttattcc ctatggatg ggcggcttc ttaccttttc 300
tcctttgtct tccgcttcat ctccatgggg gaaaatcacc attaaaggac ctcatgg 360
ctcatagatc cagcctccat agaagctcca c 391

<210> 1014
<211> 332
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1014

ngaccaatcn cgacccaaacc caggcatagt cggtcagtga gaactctgtg atgtacctaa 60
acaggcgagc tcctggcagt caacagatca aaggaacaaa gaccacaaag caaggaggct 120
tgtggggct ggccagctgt gaaaactgtat tgatatgtga gatatggct ctggtaatcg 180
attaccaagg gtgggttatac gattacaagg ttataataatg aaggaggcta acatggctc 240
tggtaatcga ttaccacggg gtgtaatcga ttaccaggct cgaaaacgag gtcatgaagc 300
catgagggtct tctggtaatc gattaccagg ct 332

<210> 1015
<211> 348
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1015

gcttctatgg agctggatct ttgagcttaa taagggcctt caatggatg tntcagccat 60
ggagttgcag cgaaagataa aggagaaaag gtgagaggag ggcggatcca ctagagaata 120

agccatggaa ggagaagctt caccaccaag agagtgcctt agataagaag ctttagagagg 180
aagcttcaat ggaggaagag aatgagagag ggagagagag agaatggtgt gaaaattgaa 240
ggagaatagg gagataagtt gaacttaaa gtgtgtctca caagttctc attcatcaa 300
agtatgacaa gtgttacaca tgtttctatt tatggcttag cacatggg 348

<210> 1016
<211> 375
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1016

cgaagaagtt ntttctttta catgcccaac tctctcgagt gacatngca ttgattggtg 60
tatTTtatgt tgcacatcttag tctctatcat atcctatgtg catcatgcat catcatgtgt 120
gagtaaggag aaaatttcta atgttataaa atttcttcag aaggcaaaac ttttggttt 180
aatccattac aaccttacag taatcaatta cacaaggttg tttaagcttg catacgat 240
tcttgcattcg atttaattaa ttacagcctt atcctaattcg attacaccaa ttgttttaag 300
acaatggttt atttatntaa tagtctatgc tttaatcaat taccatgtga tataaatcaa 360
tacttctctt tctat 375

<210> 1017
<211> 400
<212> DNA
<213> Glycine max
<400> 1017

agctataacc tcacatgtccc tcacagtctt tattattggg agccaatcca atccttgcgt 60
tcggactctc agccacttat gatagccgcc gatgatccca ttactgcttc ccctaagctc 120
tctgtccctt ctacacgccc catccatgc ctgcgaact cttggagta ccctcgcggtt 180
gtggtcacta aaacccccgtg cgatgaaagg cgtgatgctt tcgtctaatg gcgcctct 240
catgggttag ccaagctgtc ttatggtgag aacgggatta taattaatac aaccccttgt 300
tcccatcaag ggaacatttgc acatccttc gcatgaagat agaattcctga ttcttccttc 360
cttctagcga gggAACCAAT taacagacgc cccccatgc 400

<210> 1018
<211> 267
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1018

tattacaagg ctttaagtgg aaaccatgtat ttcacnctac ccttaaggaa ttntggagct 60
cttgaaaatg tttgggaata agtgggagag ggggtatgtt tcattgggtg atattgttt 120
cgtggccatg cttgatgtatg attttggcca tgcttgatgt atatacatat aatgcctata 180
tggtgctta tattttaaat gcttgcaat gctactggtc acgttcaata aaaaattaaa 240
tagaagaaga atgatgttga ataaatg 267

<210> 1019
<211> 389
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1019

gcttgcgagt ctcccttggc atttccttga catatgttct ccttatgtct aaggtctcaa 60
ggttatgaag tttgccaata aatttcagaa gctgcgttgg catcaactgaa tatcgtaagt 120
ttaaatactt gaagtgtgct agattcccc aattttcagg aacagaactc aatggactat 180
cttggaaaatc aagtacccatc aatagctgt actttgttagg gatTTTGC acaaagttgt 240
tcattaatgc tgattcttta tctgaaaaaa caaacagtga tcgggtgtgt gaattntccg 300
tactccccat taaaaatcat tggagaacgg tgctattgtat aagcgttgaa tcatcccact 360
tggcattggc tcatcttctt tactaatat 389

<210> 1020
<211> 291
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1020

cacagacaca cttagtggtt gacattaact attagtagtg gaggataata caaccagttt 60
aatgcgttgtt tggtttgacg ttggagaatt gattntgagt ctaaaattaa tattagatat 120

atttgtaaat atctggttt atgttgaaa agaattcgaa attaattcta ggtccataat 180
tgatttgga ttgaaacaat attgagtagt atctgcccta gattcaaaaa tttgtattga 240
attttatttc taacttgatt ttataattaa acattcagac ataaatcata t 291

<210> 1021
<211> 383
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1021

agcttaggttn tggcaatag caccccacct gacgtcccc aagtctcctg acccccgcga 60
catatctcca ggtaccactc tgtggtcaac gaataaaagc aggaagttca cccttctaca 120
cttcctcatc tcaagcttgt aggattatgg ggtacccatc acatgtggta ctaggtggcg 180
gtcgggcaat ggtgcacaac aagtttcca catccacaat gcgcgcataa acccaccatc 240
ccctgtgcc cacctccaac tgagctcacg tactcccacg tagcccatat cctcgttct 300
ctcaacacccg ggtccccatc aatcctccc aagttccaca acatccaagc aaaacaacat 360
tcanatagca caagctatca cag 383

<210> 1022
<211> 396
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1022

ntatcanatg gatgtaaaga gttcatttt atatggctnt attcaagagg aagtatatgt 60
agatcaacct cctagattt agaattcaga caagccta at catgtttta aataaaaaaaa 120
aaggctttat atggctaaa gcaagccct agagcttggt atgagcgtct gagtaagttc 180
cttttagaaa aggatttctc tagaggcaag gtatgata ccctttcat aaatagaaaa 240
tcacatgaca ttntactggt tcaaattttt gttgatgata ttcatttttag atctactaat 300
gaattattat gcaaggaatt ctctcatgac atgcaaagtg agtttgaatt gtcaatgatg 360
ggagaactct aattgtttct tggataaaaa tttaaac 396

<210> 1023

<211> 325
<212> DNA
<213> Glycine max

<400> 1023

agtgtttctt ttgcaagaag aaggcacaca tgaaaaagaa ttgccccggg ttccacaaat 60
ggcttgagaa gaaagggtgaa tcaatctcat tagtatgtta tgaatcta atggtagtg 120
gtaatattaa cacctggtgg attgattctg gatctactat tcataattgca aattctttac 180
agggtatgca aaacctaagg aaaccagtgg gaagtgagca aagcattta tcaggcaata 240
agctaggctc acatgtggag gccattggaa ctgcatttt gacttaagt agtggctta 300
ttttaaaatt agaaaggact tttta 325

<210> 1024
<211> 381
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1024

taggagttnt ccaaccatca agttgacctt gcttgaagac ctccatcctt catgttatac 60
ttgagagttt tccaaaata tcagacaaga tggatatgt aagggactt catttgtagt 120
acactgccat aagtatatct gtctatttgt ggagttgttc agttaccaag tagcactgtc 180
atgttaccag aactgactga tattatagct ntggatgga aaggtagca atggctaaat 240
cacgaagatg gtgaagaaaa agcgggtca atagtatctt caaaggtaga acggctntgt 300
gcctaattgat gcaaccattt attaattttt canaacggtt catgcacgtt gttcatgtga 360
aagattttaga ctgattgata t 381

<210> 1025
<211> 289
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1025

gcttatctt gttntacaa ccaaagtcca tgtgaacctt gagtaatcat ctactataac 60
taagccatag taatttccac ctaaactcag ttctagaggg accaaataaa tcaatgtgaa 120

aaagttcaag gggtttgaa gtagaaacaa cattttact ttgaaaggag ttttaactt 180
gcttccttt nttacaagct tcacacaatt tattttctc aaacttaagt tntggaagac 240
caattactaa gtcttccta actagatgt taagatgtg catatttat 289

<210> 1026
<211> 410
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1026

ctcaagctta tgattggatn gataaatgtg aacgtgagag acctctgagt gatgcagaga 60
gtagctgacg aagagatatc gataggtgag gggagttat tttggttta tagctttga 120
taccatactt gngtttagta aacccaacta tggggatgta tgattctccc tatgcatgct 180
agtttcaag aaaactgtgt ttntgactaa tggatgcga tatatttctt attgatgaat 240
gaaattgtga atgatattgt tgtttatga aacttgtgtt gtctgaagac ctgtgaaatg 300
tgaatccctan gcatgaaatt atatgtatat atgtggaatg cgattactga tcatgttaat 360
attgatgata atattgatat gaaatgatgt tgatattgag atgagatgt 410

<210> 1027
<211> 360
<212> DNA
<213> Glycine max

<400> 1027

aacaaaactt gtgttattca tcttttcat tctttctcc cttgc当地 aagaatttg 60
caaggactaa ccacctgaat tcttttgtg tctctttctt ccctttcca aaagaacaaa 120
ggactaaccg cctgaattct tttgtgtctc cttctccct tgtcaaagaa ttcaaaacga 180
cacagtctga gaattctttt gattctccc tttccataa acaaaagatt tcaaaggact 240
aacccgcatga gatatctttt gttccctt cacaagttt caaaggacta accgcctgag 300
aactttgtct taacacattt gagggtacat cccttgtgaa caagtagagg acatctactt 360

<210> 1028
<211> 409
<212> DNA
<213> Glycine max

<223> unsure at all n locations
 <400> 1028

tctatggagg ctggatctt gagcttaat gatgtnttc aatggtgatt ntccaccata 60
 gagttgcagc ggaagataaa ggagaagagg tgagaggaga cgccatccac tatgaaataa 120
 gccatggaag gagaagcttc accaccaaga caatgtctt gataagaagc tttagataagga 180
 agtttcaatg gaggaagaga atgagagaga gaaagtggca tggaaaattg aaggaagaaa 240
 gggagagaag tttaactntg aagtgtgtct cacaagactc tcattcatca aagttgtgac 300
 aagtgttaca catgtttcta tntatagcct angtcactaa catttcacgt gaatctaaga 360
 ggaatattcc aagaatatcc canatgcac ttaacatatt ccaagaata 409

<210> 1029
 <211> 521
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 1029

cttcgttcc gtctatngcg ngcccgcat ccttagagnnc gcctgctgca tgcaagcttg 60
 angaganaag gngaatattc tttttcttct tggatgaccc aangtggcaa cgtgcttcat 120
 ttagttaaat atcgtacaca ggctcattca ttgtggacgg taacccagcg gggattctta 180
 agacaatgat gactcacacc ttcaattcgt tggaaagctct tctaacctaa atttgacata 240
 ctacactgga agagcactta ttcatttgca ccgangggtg gcatatgctg ccatgatgaa 300
 tcgaacatct aggaaaccat gcccaactca cagaatttaa tatgcggcac actactaagc 360
 ccaacgtgaa ttgcacagga aatcatggtt gcgcgttta gaattgactt catcacaagg 420
 gtagatacga taagacatgt actgtgtttg atacacctca agctccaatt gatcgccgag 480
 tcatctagga gatgatcgtg aatgcagaca caacctcacc t 521

<210> 1030
 <211> 87
 <212> DNA
 <213> Glycine max

<400> 1030

actttcaggc gactttctct aagcgccgca ccggaatctt caagaacgca agtgagcttg 60

ccaccctctg cgacgtggac cttgctg

87

<210> 1031
<211> 460
<212> DNA
<213> Glycine max

<400> 1031

aagcacctga gctgcagcta tgctgcata ttacaataga ctttcttac ctcagcagca 60
aaatcaacca caacagcaca attatgacct cttagcaac agatacaacc ctggatggag 120
gaatcaccct aatctcagat ggtctagccc tcagcaacaa caacagtgc ctgctccttc 180
cttccaaaat gttgctagcc caagcaaacc atacattcct ccaccaatcc aacaacagca 240
acagccccag aaacagccaa cagttgagac ccctccacaa cttccctca gaagaacttg 300
tgaggcaaat gactatgcag aacatgcagt gtcaacaaga gaccagagcc ttcattcaga 360
gcttaaccaa tcagatggga caattggcta cacaattgaa tcaacaacag tgccaaaatt 420
ctgacaagct gccttcttaa gctgtccaaa atccaaaaaa 460

<210> 1032
<211> 419
<212> DNA
<213> Glycine max

<400> 1032

acaggcctat atgacatctc ggactatgat taactccctc taaccccaa gtaccagcaa 60
atccagaggt aactctacaa actctcaaag catcactctt tattcactcat agcactacat 120
tctcaactatc taaccctagg ttaactctac cctacatctc tagcagattt ccataagcaa 180
ttgcaaaaca cagacatcac atgcatcatc atagacactt ctaaaccaga acggaaagc 240
gtgactcaca cctgacatga cgaagttaac atgtttcagt gagattctga cagataccat 300
ccagaacata aaccttagttg actacccatg atattccaa aacaatccca cagaatatgt 360
gagaagatgc taccaacctg aaattgaagt cccactatag ggcgcttacg actccgaaa 419

<210> 1033
<211> 448
<212> DNA
<213> Glycine max

<400> 1033

agctcgaact tgaaataggg tcaggattga tcttacgtt cctcatggct cttgaagaaa 60
taaagagaat gaaaagtaca aataaatatt attttatttg taaggattaa aaatacattt 120
aacacctaaaa attaacacgg attatgacta tttttataa aattatattg ttgtttttt 180
ttaatttta gataacttat gattgcaata ttggtaatg aatataaact tcacccaaag 240
ttaatagttg attttatata attaaaacca aaagttaata ttttataata tatatatata 300
tatatatata tatatatata tatttcattt cagtaagaaa aaatcatatt atatataaaa 360
aaagctctat ataaaatttgc atagagggtc tcaactctca agcaccttaa gtcagcctac 420
tagggtcatc acaggtaccc gaagataa 448

<210> 1034
<211> 356
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1034

cggcagttnt tacgtatact ttggaagtat atatattaac atcacgcact tatacgag 60
taagagcagc tcttctacgt ggtcattgaa ggaagattt agtgttgta ttttttgta 120
acttacataa tggctctg atgatattcc gtgaagaatg ttaacgactt ctcgttttag 180
ccatcctatt ggtctttca ttgcgtac atatatatga attatactta ttgcgtgaga 240
atattctta cttaacttgag actattctaa ttgtgtgcc tttaccagaa tatcttcacg 300
ctacactgga atgacaatgg tggatgcctc tctatgagaa ataatgacca catgtg 356

<210> 1035
<211> 449
<212> DNA
<213> Glycine max

<400> 1035

agcttgcgtat ggtatataata tggatgcctc tctatgagaa ataatgacca catgtg 60
taaatatattt gtgtatacta aaagcaaatg cttaacatta tacgtcttc gtactcaagg 120
atataccaac attgaaggc ggactgtctt gatgtagcga ttttaacagc gatgacacta 180

tctttgttct atatattaac tgtcatggag atggagctgt ctttgtaaac aatctttgtc 240
attatccatgct gaaagatgaa catgttgttc gataggaaaa taatgaaata ccttatataa 300
atgtattgt aagtttaag aataaactgc tcaaattgcc agcagttgt cgtagaatag 360
tagtgcaaat tcatgtatgt aatgtattag aatacagaat aaacctatgg tgccgttatt 420
aattgttatg agaacatctt gggcttgc 449

<210> 1036
<211> 82
<212> DNA
<213> Glycine max

<400> 1036

tagcacatgt tgggtccaa tgaactcttc ttgacagagc atgtgtgaa caggaactct 60
tagaatgatg tgttagaatga at 82

<210> 1037
<211> 264
<212> DNA
<213> Glycine max

<400> 1037

gagcaacgca cagctcacat tctctgttcc aactcctaaa caattttaa ttgggtggctt 60
ctggatttat gtcaatacat caaaatctt tggataactt gtgtcatcat gtaatgcttc 120
ctctactact gattcgataa aacagataaa aaaacactaa aaaatgaaac ctaatatcat 180
caacgacata aagcataaat tctagtatta gtatcacca aagtttggc tgctggttt 240
gtgcccattc ctcacattt atct 264

<210> 1038
<211> 429
<212> DNA
<213> Glycine max

<400> 1038

agcttctact tatgtggcag ggccggcttc ctttcccttc ttgtctccaa cgcgaaacttt 60
gaccattgtt ctcccttccc gcgatgcttc tttcatgtc tgcctgagtg ggcttatagc 120
ctaaaccata cttccccacga ttaccttggg tatttatcag tctagttatg ccggccgttgt 180

| | |
|--|-----|
| tttttcctaa acccataaccg ggctcataac cgttcccaa cataactcg | 240 |
| ccgctgcatac ggacagacta tgctgccaa agagggagtc cacggaggaa atgctgacca | 300 |
| cctcaaaaga ctggaaagta gtttctaacf attttctgc ggctccaca taaggcatgg | 360 |
| aggatgggca gcttaccaag atatcttcct cgccctgacac gatgaccaag tgcccctcta | 420 |
| ctacgaatt | 429 |

| | |
|-------|-------------|
| <210> | 1039 |
| <211> | 437 |
| <212> | DNA |
| <213> | Glycine max |
| <400> | 1039 |

agcttcgaag ggatgaaagg gatgtttga gggtgatgag tatgatatta gaggataagt 60
tgaaggcttgc aaaggcat aatagagttt gaccgagcag tttagttata cagacgagaa 120
catcttgaca atcattgatg ggtataaaga agagctaagc ctatcgcca gtcatacggtt 180
gagactagag gatgagcacg cgaaggtagt ggctctgcaa gcggaaaggaa aagcaagaga 240
gagcgtgata taatcattgc acagggaaagc cgtaaaatgg atggatagat tcactctcac 300
tctgaatagg agtcaagaac ttccaaaggct tttagccaga gccaaaggaaa tggcgatgt 360
gtacccagct ctcgaggaag ttcatggct tctcgattat tgccaaacaca tgttcgaata 420
gatgtgcctc ataatta 437

| | |
|-------|--------------|
| <210> | 1040 |
| <211> | 408 |
| <212> | DNA |
| <213> | Glycine max |
| <223> | unsure at al |
| <400> | 1040 |

tcatgatgat gaaccaagca attttgatga tgccaaaagc ccttgtgatt gattcacgac 60
ttcaagatca agcatcaaga atccaatcca agattcaaga ttcaagagaa gaaatcaaga 120
agcaacaagt caagacttca cacatgataa gtattaaaag atttttcaa aaaccaaata 180
gtatagttnt gtttacaaa agaattttct cacattgtct aagttaccaa agtgattact 240
ctctggtaat cgattaccag ttatcagtaa tcgattacca gtgaccaact tggttccaa 300
aatgtttca aatggttgc aacgttccaa aatqatttc aacagtgaat cgatacacta 360

tatattaatc gttacagtga atctgacgtt gaatcaatct attggaga

408

<210> 1041

<211> 423

<212> DNA

<213> Glycine max

<400> 1041

agcttcgaa gcatctacaa gatgtattaa ctgtgttagt tgtcagccat gattagagg 60

tgcttcaata tactctatta gttttgttt atctggctat tcagtcatta ttgatgctca 120

tttatcgta tatgtatggg tttgtcttc tcgtgattga tgatttgcga ataccattag 180

aaattcta ac ttcgattgtt acgaaagact gatttaatgt gagcttcaa attgtgctga 240

tatatacatg gataaacaca tgacgttgat acttataatga ctgtggtgct tcaatgtata 300

cacttggttt tatgttaggca acaacttcta gaatgtggag gctatgagct cgtgcacatc 360

atgaactcaa taatcactta tgtgataatg tatggctgga agtcagtaaa cggaatctt 420

agc

423

<210> 1042

<211> 437

<212> DNA

<213> Glycine max

<400> 1042

tctcgccatt gacaatggcg gtacgcgtat ctcgccagta cttctggcga catccatgg 60

aaaacagacc ccctctgtaa atacttataa aagagacccc tttacgtaaa tagtttgtaa 120

aggtgaccct ctacagtaaa tttaccactt taaaataacg tttggacatt ggattttcat 180

ataagttatg gatgcttata taaaatagtgt gttggaagca atgcttgata acccatttt 240

atgcatctaa gttaagttac aagaatatac ctattacttc cttggacgca tgtttagctt 300

gcgtagttca ataaaactata tcaattcacc aaagtttaat atctcttatt atcattagaa 360

ctatgaaatc ttacgggct tcacattatt catgttttg gttccggcaa aatgtttga 420

ccaacttgcc atgtaag

437

<210> 1043

<211> 407

<212> DNA
<213> Glycine max

<400> 1043

taacttgagc atctctgact atgaatcatg caacataata gggAACAAAT tcatagtAAC 60
cctcgagtac aagaccgaga tgactataga gtatgtatccc ctcataatttG atgtccccat 120
tagcaacaga cttagcatat gttttatTTT catttgaaaa ttttgtgaca tgtgtcgac 180
taaatccaac gaataaacac aaaatacatg ttataacaAG gattctgtGA taaaattatG 240
tgcacccTAG gacgttaATTc tataacatgt tcttagttGA gtacgaggct ttacacCTT 300
tgcttgacaa tggcaaAGGA gatgcacata tagagtaACT agcgagctat ctaaataACC 360
tctcgtgaca tttagaccAGC ggaaAGAAAC atgatgacCA aactctG 407

<210> 1044
<211> 325
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1044

tTCGGCCGAC actggcgtGT tcccatgcAC tccggcGAGA acacattGA cCACCTCGAT 60
cagataaaaa gacattaACC accggTCTTG atcggaaaaA atgctggTTG acgtcgGCCA 120
tgatagatGA ccgatcgagg tctgaaaATA aaagaatCAC cgatgacGC cgatcgAGCA 180
tatcctaatt gacatcatCC aaatattATC cagggattGG atagaaaaAA caatagCTGA 240
taccagTCGT tatgttagtCC cgactgacAT ttgtcAGCCG acattgcACA gtatttNTT 300
caaacgCTGG ccgataatat atctt 325

<210> 1045
<211> 344
<212> DNA
<213> Glycine max

<400> 1045

gctttacgga cctatcaaAC tcAGCTAGGA ggatTTGGA gACCCGCTGT tgCGATAAC 60
gaggatATGG gCTACGTGCG agtacgtGAG cttaattGGA ggtgggcaAC acgagatGGT 120
gggtttatGC gcgcattGTG gatgtggAAA acttGTCGTG caccatCTTC cgaccGCCAT 180

ctagtaccac atgtgattgg taccgcataa tcctacaatc ttgtgatgat gaattgttga 240

aagaggagac ttccctgcttt attgttgacc acatagtggt acctggagat ctgttgcga 300

ggtaggaga ccatggggac gtcatgtggg gtgctattgc ccaa 344

<210> 1046

<211> 420

<212> DNA

<213> Glycine max

<400> 1046

taatgacatt gattatgaca tcacacgact tgctatTTT agttcattt tttccaaga 60

aatatcggt acctttcgta aaagaattct gtttgcgtcc ttttgcagg aaaaaaaaaa 120

aaagagattc tgattgaatt ttagtaact atttctaaa ataataattat tatgagtgac 180

aactttttc ttatcttaac actctgttt gctgtatatt aagactctga ctcaaaatca 240

tcagacttgt ctataaaata agtatcttct acttccatcc cagtaaaaat cccacatgaa 300

ggacttaaca aagcttagatt actttgtgac tcttataat ataactaaga tgattattac 360

gaacacatgc tatgggtta tcttcgaaaa ggaacacatg aaataactcg attttaattt 420

<210> 1047

<211> 376

<212> DNA

<213> Glycine max

<400> 1047

agttggaagc aaaacaaggg agcaagctt gggAACACTT tcttcaagaa ctaaaacaaa 60

gtcctttaac ctTTTccatt ttcattcctt ttactatcct catgtatttc tggattggat 120

tcttctcctt gcatcagtag ttctacaaaa tagaaggagc agaacacaag gaaatttagat 180

tatTTAGGAT agtcacatat ctcctatgtg ttAAATTAA gtagctgaa ccattacgta 240

ctaactctga tcactctata tgtctattgc ttgctatATG attcgctaac acttttgaca 300

agGAACtGGA tgagatgaag cacataacta ctggAAATTG tcgcaagttt ttattacatc 360

attgttagttt atgttt 376

<210> 1048

<211> 448

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 1048

agctgtgctt ctacactacc agcaacaata ttttatcaaa gagaanaact ctagatgagg 60
gttcactgtt atcaagcaag tcggagaccc agcatgacca caaattcacc tccactcctt 120
atgttcccat ggacctgggt atagggcccc ttttcaactc accgtgtgta caaatagtgt 180
tggtgtttat gtgcataaaa tgaataaata tctatctcat gcttacattt caaaagcaca 240
ctaaaagcaa aaaagagtta tatacaagaa cgtaaaggaa ataaaaggaa accgacaaaa 300
gaggaagtca tgatattgca cgagattaga aggcctaact ctctaaaaac agtccccagt 360
ggagtcgcca aatgtcgcaa cctacccttc ggcgggaggg cgacgcggng ctcacgggtg 420
tgtctccaa gggaggaagg ctcacgga 448

<210> 1049

<211> 447

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 1049

tgtaggcctt gaatcttctt catcaatgga gtccttgct tcttgaagat aaattggaag 60
cggaatggag aaggaggaaa ggtgattaga gatgccactt caaggagaag atgagtcaag 120
aacaagttca ccgccatatg aagatatgga tataagagct taaaggtacg agaagatgag 180
tggagggaga gggagagaaa gggcacgaca tttatgcctc agatgaggta taaaatgtga 240
agtgttaattt ctcnatgtat caaagttaaa aatatgcaca cacaaggcct ctattnatag 300
tttaagtgtc atacaaaatt ggagggaaat ctgaatttctt attcaaaattt cacttgaattt 360
tgaatttgtg gagccaaatt tggagccaaa attcactaa ttaggattgc atcatccctt 420
ccnctctgaa aatgaattga cctcaaa 447

<210> 1050

<211> 448

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 1050

agcttgc aatcaaata ctcctacgtc tcatactctag catgcatttt ctttctttac 60
ccactcctca cgtgtggttt tttagggaaa aaacaccata actaaacgca cgcacatggg 120
tccctatcg accagatcca aatctagaac gatgggtat caagaggaga cacaggaaca 180
gatgaaggcc gacatgtcg ctctgaaaga acaaataggcc tccatgtatgg aggccatgtt 240
aggtatgaag catatcatgg agaagaacgc ggccaccgc gcccgtgtca gttcggctgc 300
cgaagcagac ccgactctct tagcaactac gcaccaacct ccctcaaaca tagtatgacg 360
gngaaggac acactgnngc acgatggcag ccctcacctg tgatacaacc gagcggctta 420
cccttatgga ttgccgcccc actattca 448

<210> 1051
<211> 399
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1051

ctgagggaaag aagccacaag aacacttata tgaaggtcg atgtataaga agaacatgt 60
ttgctctccc tcccttgaag aactcgatca caacaatggaa gaataaaggat tctaagttt 120
ttttttcttg gagaagcgag gacatataag gctttatgct tgcttcaaata gaaacttggt 180
tacgcttaat gttgacaaga tcaaactgtat gacatgataat atcattgtat agccataatg 240
ctgccccata tgcattttctt gccttttgc tntttacta gaaatgacta aagtcgactt 300
aagcaaaaat ggtaaaaact ctttctgtaa aactgaaaac cttatctaataat ctttagatag 360
tgtgctacat cttggatata gtgactcata ggaacttgc 399

<210> 1052
<211> 448
<212> DNA
<213> Glycine max

<400> 1052

agcttctccc tataacagct tcaaaattctt atattcagca cactactgtat attcaattc 60
actcatgtta ggtggcatcc ctaaattggg aatctatagc ctggccatgc ttaaatgctt 120
ataattttctt agatgttcca gcttaaatgtt tttgactgtt ctcagcatga gaatttactt 180

ctcacggtgc tcaaacaagt gtgttggta cagctataaa gatctgagtc aatgaggttg 240
ttaggcaagc ttgtctctgg gccctctctt taaattcctt gttcgtcca tttgggctcc 300
cttctacttt atttgttaat gattaatggt gatgggtatt tatatatcca tttggaaaat 360
gactaataga gtcttggtt tagagttcaa ctgaacataa cctactttt ttaaaaacaat 420
aaaatctacc agtgttattt cttctcta 448

<210> 1053
<211> 440
<212> DNA
<213> Glycine max

<400> 1053

tcatcaagga tcttggtaa atcatccat tgttcagtgg ttgttttta ctctgtcatc 60
ttgaagggtgt acagtttttgc ttcaagcat agccaatttgc caaggactt tgtcatatac 120
aatgactcca gttcaacca cattgagggtt gttgtctttt ctcttgcac ttctcttaaa 180
gctttatctc caaagcatag aatgattgca ctgctggctc tatcaatcat ctctgatttc 240
tcctttgagc ttagagattt agacatccctt tcttctcctt taagagcttc tgcacagcc 300
tgttgaatca agattgcttc catcttgact ctccataacc cgaagtcatt gtcccctgaa 360
aacttctcaa tatcgttattt tgatgatccc atcttcttg gtcttgatct tgtccccata 420
gacggcgcca cttgttgatt 440

<210> 1054
<211> 447
<212> DNA
<213> Glycine max

<400> 1054

agcttctacc ccattttcct ataaataggg ggagaagtga atggtaaaaa tgttcagccc 60
tcctggtaat tcgagaatca cttgaaatattt gcggaaaaaaa ttgtttccgt gaagaaaatc 120
caagtcgatg cgcttccgtt acgtttccgtt gggtgatttc gcaaagattt tcaaccgttc 180
ttcgacgttc ttcggttcatg cttcgccgtt cttcggtctt cagccggtaa gtttccgaaa 240
tcaaactttt caatttcatc tatgtacgct tagtggtcctt catttggttt cacgtgcttc 300
tattatcatt tcatttactg tccgtacccc ctttgacgt gctttattca tttgcttaag 360

tcattttgtc gcctaataca atactaacat aaatttccac tgatcgcttg aattgtaata 420
tccgataatt tctgttaaaa tgaaatc 447

<210> 1055
<211> 446
<212> DNA
<213> Glycine max

<400> 1055

gctctctaag tgaaatcagg tgcagccatc tccctaagag tcctctcaaa aggtggaggt 60
tgagccatgt tctcagtatg aaaattägt a t c g a a t g c t a a t t c a g a a 120
tcaccatcaa cataataactc agaatgctta aaatgctcaa aatgcacaga atgatcagga 180
tgcacactat gcctaagtaa tccatgaaag gtttatctt tttcaggaag gtttctaaat 240
cacctggatt gcccttagtc atgcattata tgcagcaa atcatgtttc tcaaacaagg 300
accagttggag gtttaaaact acaactatag tcaa atgata tccaa atgag ctgaaatttt 360
atgagtaaca ccctaaaatc atgaaaagat agaaca aaaa tttgcagact aaaattcact 420
aactatgaaa actgactaaa gaaagt 446

<210> 1056
<211> 204
<212> DNA
<213> Glycine max

<400> 1056

tgctggtgg a g t t c g a t g t a t g c t a a t c t t t g a g c t t c a a t g g t g 60
agtgttccacc atggacacgc cacggaaagt cataagataa gaggataggg gaggcacctt 120
tcactatgga ataatccaag gaagaaggag cttcaccacc aataattgcc ttggataaaa 180
aacttgcaca agattcttc ctgg 204

<210> 1057
<211> 369
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1057

agcttctagt cgtcataga cttctctng ggtctgacta tcaa acgttg catctgtgca 60

ttcatcgcat tcactaacag acgttggcg ccgtccaact gatggtaactc gtgaccacca 120
ccacctgcta cagccataat ttaacaggaa aaaaaaatgt gcaataaaaa ctattaaggt 180
ttcaagacct cacaacactc tactcacgtc tgtagatgt gagtacactc gtgttaacg 240
ctctcaatat gctatcgta aatgtattcc ctctgcctt ttaccactcg agtggactct 300
taagttcctg gatggaccaa attacacaca catggtaata tttaatcaga ggagagacta 360
tatgatgat 369

<210> 1058
<211> 441
<212> DNA
<213> Glycine max

<400> 1058

agcttgccctc aaagagggtcc aggattgata atggtgccga aggaactagt tccgctcccg 60
agtatgacag tcaccgcttt aggagcgctg tacatcagca gcgcctctaa gccatcaagg 120
gatggtcatt tctccggag cgacgcgtcc agtcaggga cgacgagtat actgatttcc 180
aggagaaaat aggtcgccgg cggtggcat cactagttac cccatggcc aaatttgc 240
cagacatagt cctcaaattt tatgtcaatg ctggccaac agaggaggc gtgcgtgaca 300
tgaggtcctg tgtgaggggt gagttatcc tgtagatgc agatgctatc ggccagctcc 360
tggatatacc gtttgtttt gaagaggccc aggagtgtga gtatggccag aggagaatc 420
ggtctgatgg gttcgatgag g 441

<210> 1059
<211> 407
<212> DNA
<213> Glycine max

<400> 1059

cgcataata atactggac agtctcaaacc cctgatgtat cagttcaga tccgtggagc 60
caatgcgcag tggacaagt agagtcgcta aaatcattgg tcagactcct acctatgtgt 120
gcctcggcgc tcttgcgtat ggcgtccaa ggctcattct ctaccctgca agcaactacc 180
ttggaccgaa agctatttg caatttcaag atgcctgcag ggtccttcaa tcttacatg 240
atattgaccc tatcaataga cattcccttg tatgaccgca taatggtacc tctactagcc 300

aaatacaggg gcttgccgaa tggattctgt agtaaaaactc caattggat tggattgctg 360
tttgtatgcg cagctaaagg aacatcagct gttagttaaaa ctattag 407

<210> 1060
<211> 430
<212> DNA
<213> Glycine max

<400> 1060

gttattcgcg catatttgc tcgggtcgct ccatcatacg atccatgaca cgccatgcat 60
cctatctgct gaaaaacaca aaatgcttag cgtactaatac accgtagctt gttaacatga 120
acgtattaaat aaatcttagta ctgcgctcac tcacctatga ttccggccct gagaagaaaa 180
tgaatctgga aaatgagaag gcaacaacaa cagcgcgtga cgtaaactct tatgataagg 240
ggagagaaaat gagattagac gcttacgcta tatagaacga tgcattgccc ttcttttagag 300
atgacgtgac acacttaggtg acctcttttg caaaactaaa tttggggccc ttgtactagg 360
tactataacct tacaacgggt ttccggtgta atgtttctta agatttacag agagttatac 420
atattgttct 430

<210> 1061
<211> 380
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1061

atcaattcat attggatgca ctccaacttg taatgatgcc ctattcctt cctcagagac 60
tgatgccatg acaccgtcac ccttttttc agaattcagt gaaaacccct ctctcttgt 120
acaactttcg cgccttataa cgacattcaa atgttctaaa aattaattgg agagtgaaga 180
acaaacctca ctacgtgata attgtcccc catccggcc ttgtgcaaac attcatgtga 240
taagattttg aaagtggaa tagtagatcg aggccaccta aatggaaatt gaaaaaaggg 300
tgtcaggtac aattatcaca cactntttta caatgatata ttgcttagaa ttcaaaacat 360
tcctaattgtc cacaacaccg 380

<210> 1062

<211> 439
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1062

agctntgttt catagataca tatatacaac taatataaaaa tgtaacaaaa atgatgaatg 60
aagataaaaa aaaatattgg aatgaaatgg ctatgacaat gatgaagatt aacaaataac 120
cacacgaaat attcaacaca attcatatta agatcagg tataatttc aagactattc 180
tcaaataaac atgacatgat ataacacatc agtattgctt atattatgg actttaatg 240
tagccttatg gtgaagtaag tagtccttt gctgtaaat ggaaggatta actagaagac 300
acatggcattc tgtagatag taatgacagc atgcactatg ttatatgcaa caaagatctg 360
gtgaacccaa ctatactggc tggatggaca aaactcgag atttctatgg actcacagga 420
tatcatcaag tgaccatga 439

<210> 1063
<211> 410
<212> DNA
<213> Glycine max
<400> 1063

ctatatttc agtagatgaa tatgaatccg cggccaccc atgtactcct ctaaggacaa 60
tagcatcatt tgtagtgcactg aattgttagg agttgaaagc catcttctca atcaaactcc 120
tagcctcagc acgggtcata tcaccaagag ctccccact agcagcatta atcatactcc 180
tctccatgtt gctaagtccc tcatagaaat attgaggaag gagttgctca gaaatctggc 240
ggtgaggca gcttgcacac aatttcttga atctttacca gtactcatac gagctctctc 300
cactaagatg cctaattgcct gaaatgtctt ttctgtatggc agtggctcta catgcaagga 360
ataatttctg caagaacact cttAACGTCG tccaagctga aaatggacct 410

<210> 1064
<211> 432
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 1064

agctatgcgg atttggctt cgccagtgaa atgttcgaag tggatctgan aagaggcaaa 60
tttaatcatc ctgcttagac gaatgagaaa actgcggcaa ataaagaggg tgaggatgag 120
ggagaaaaccc atgctgtgac tgccattcct atacggccaa gtttcccacc aaacccaaca 180
atgtcattac tcagtcaata acaaaccacc tccttaccca ccacccagtt atccacaaag 240
gccatcccta aatcaaccac aaagcctgtc taccgcactt ccaatgacga agaccacctt 300
tagcacaaac cataaaaaac accaaccaag aaatgaattt tgcagcgaaa agcctgtatg 360
attcacccca tattccggtg tcatatgcta acttgctccc atatctactt gataacgcaa 420
tggtagccat aa 432

<210> 1065
<211> 422
<212> DNA
<213> Glycine max

<400> 1065

tctatataat ctgaaccatt ctatcaataa acacacgtcg agttgtattc agaatattag 60
agtttatctc ttattatctt agtgagagtg attctcctaa attcttgagt gattcaagaa 120
caccttggct gtatcaaagg actttcacaa cctttgtgtg ttgcctcgc tggaaagagt 180
gattcttctt ttactttcat catcacccctt gttcttcaa atcacaattc cagaagatcc 240
acctctgccc agagatatct cgtggccata acttccattt tacgcactca aattaagaga 300
ttcttgagcc tatattgaat ttcaaaacga gacccttcac ctcgttatgg aatcacctca 360
tttggagccc tgttagcttca gtattgccccat ttcttatattt ctgtccagcc accacttaac 420

ct 422

<210> 1066
<211> 435
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1066

agttcataa atagactgaa tcaaggagaa aattatcgac gacccccana acataaggta 60
tttaacacca cttgtcacgt tgttgactag ttaatagagt ttattatgtt tgcatgaaag 120
aacatgtttc taaatgtatt aaaaagcttgc cctggaaaa atcaagttat gagttatggac 180

tagagatgag atgcaaactg caatcaacat agggtacaat aacaaggctt gttctagatt 240
ttttttttt ggaagttgaa taaagttaa tctagttgtc tttttacagg tatgctttag 300
gtttacttag ataagacatg aaacagaaaag tgaccactct tgattaacta gatattgagg 360
cccttgtaaa aggtactaga ctttacaaag ttttcgcatttaatttct ctaaatacta 420
tagatattt ataca 435

<210> 1067
<211> 428
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1067

ntgatctagt gtaaaattaa ctccctan aagataaaac attcacaggt ccagatgatt 60
accgaatttg taagaacaac aactaagtca tcttggaaagt actcaaacag ataatagtgc 120
actatacgctt ccctcagaga cagaggccat gaaaccttca ccctttctc caaaattcag 180
tggaaaatca gcattcaaaa gtggaaaactt ttggcatggt aaaggacatt aaaatgttaa 240
aaaaatttaat tggtaagga agaacaacc tcactagtga taattgttcc ccaatccggg 300
cctttgcaa acattcagct gataagattt tggaaagtggg aatagtagat tgaggccacc 360
taaatggaaa ttggaaaaag ggtgtcaggt acaattatca gacactnttt tacaatgata 420
tattgctt 428

<210> 1068
<211> 442
<212> DNA
<213> Glycine max
<400> 1068

agcatttaca atgattaaga tatactcttt caagttttt ggccataaat tagtctggga 60
tctcaatcaa gtctcgagac tcttggaaagt caatggtctt taaactcacg aggttctgt 120
aaaaaaaaataa aaatacatca cattgaccat aaatcattaa aataaaattc ttgtataaca 180
ttgaagctca tcattcctagg tgtggcttctt gtgattgtga atgagttca ttgcagtaag 240
caaaagacta gttaccactt tgtcttacga ttctctaccc tacctttgca aatagaaaga 300

aaaatgtagt atgtatattc acaatcatac ctgaaccca tcccagagct tttttagctt 360
gctacgaggc atggaaatct ctacaagtcg tttagcgag aagttatacg gcaaagactc 420
aagacaacat tcataccat ga 442

<210> 1069
<211> 450
<212> DNA
<213> Glycine max

<400> 1069

gagcttatg gaggctggat ctttgagctt caatgaggc cttcaatggt gatTTccac 60
catggagatg cagtagaagg caaaggagaa gaggagagga gaggcaccat ccactatgga 120
ataagccatg gaagaaggag catcaccacc aagaatgtgc catggataag aagcttgaag 180
atgatgctt aatggaggaa aagaaagaga gaagggggaa gcacgaaatt gaaggaataa 240
aagagggaga gaagtggAAC tttgaagtat gtctcataag acgtccattc atcaaagtta 300
caacaagtgt tacacatgct tctattata gactaggtag cttccttgag aagcttctt 360
gagaaaactt ctttgagaag cttcttgag aaaacttcct tgagaagcta gagcttaact 420
acacacacac ttcttaataac taagctcacc 450

<210> 1070
<211> 448
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1070

agttctcnN cctatttgct ataaataggg ggatatgtga agataaaaag ggTCAGCCC 60
cttaggcact tctctctctc tcgaaattgc tgaggaaaat tattccgtg aagaagaaaa 120
gggttcagcc ccttaggcac ttctctctc ctgcggatttgc tgaggaaaaa ttattccgt 180
gaagaaaatc caagcagagg cgcttcgtt acgtttccgtt gagtaattac gcgaagatTC 240
tcgaccgttc ttcaagattc atcggtcggtt ctgcgttttgc ttcaagtcttc aacgggttaag 300
tacctcaaac cgagcttttgc aattcattct atgtaccgtt ggtggccac atnttgcgtt 360
atgtatTTNT attctcgTTT tcatttgctt tttatacccc ctnttgacgt gcttaagcc 420
tttatntaag tcatttctcg cttaatct 448

<210> 1071
<211> 107
<212> DNA
<213> Glycine max

<400> 1071

tgctgactgc aatcatcttc gggtaacattt acaaggctgg tagtagaaat cttgtaaatg 60
gaagtcgagt caaccttgat gcagatgact ttatacacgc atgcttc 107

<210> 1072
<211> 384
<212> DNA
<213> Glycine max

<400> 1072

tggatctgac tgccctgctt tatgtgatat acataattgg gtataaagac gactaggcg 60
cagtaactga cgtttgtttt tataatccaaa ttgtataacc tgtgataaat cattaaagc 120
cattggtcct ttattattca ttatatgatc aattctttt ccacttctag attgtgagta 180
tcaatgatta ttcatgatca caaagcatat gccaaactca gatgatagta tatcaaatta 240
ctatacttct gatcatgagg gcgagccctg gtgcattcggt aaagatgtgc ctcggtgact 300
tggggcat gggtaaat ccagaaacag cctcttgca tatgcatggg taatgctgcf 360
tacaacatcc ctacccata cctt 384

<210> 1073
<211> 318
<212> DNA
<213> Glycine max

<400> 1073

agctcctgcc tcagttata ctcatccgtg gctgtgtat cagcttttag aagatggcca 60
cctaggattc ttgtgtgcat tttccacacg caagagggtt aaccgaaaca catccagcat 120
ctaagtcggg aatcgaaattt catgaaataa tggccgacat tctcaaattgg gtctatacta 180
gattgccttg gatgagttga aactgtcggt gcattatctg aaaatgtgcg tgccctgtaa 240
aggaccatat ggataaagga gctgaccacg ccatacttat acagatgtac ttgtgctacc 300
tactttcccc tgattacc 318

<210> 1074
<211> 356
<212> DNA
<213> Glycine max

<400> 1074

agcttgagg ctggaagcag tgctttact gtttctggat gctatctaca ataaaaaagaaa 60
cccatcagtt tattagacca gaagtttatta agattaaaac agaaaataaa aacgaaaattt 120
ggcgatgtgc gcttagcgag atgcagctag cttagcacgc cttagtaaaa acaacacacc 180
ggcttagcgc aatatggttt cgcttagcca gtcatgacaa agaaattttc tctgcataat 240
tggctttgcg agcagtgcata gcttagcctt atgcatgccg caacgaatag tgcttagccc 300
atggggatgg cacttatccc gagcaacact tccaaaaatt tgactatgta atctgg 356

<210> 1075
<211> 252
<212> DNA
<213> Glycine max

<400> 1075

tgcttagctag ggttaatctc aaagcttcat aagcatatcc ttgtcagagg actgtcttta 60
gtcatcatat caggatgatt ctctatgtga agcttgcctt aaaggaaac aagtgaaaag 120
ccattgttat aaccaaaaaa taccgttgc acttctacgc ccttaacgcc tatgtggccc 180
aactataact acatccctct ctggacacat atatggttt gtcatcgagg actattacac 240
cagatggaca tg 252

<210> 1076
<211> 412
<212> DNA
<213> Glycine max

<400> 1076

gttgcttcct ccagaaggca tcgccttctg gggaaactacc tggaaggccc tagtggccct 60
ggtttctatt tgccacccctt atttactaaa tacacccctt taccttttt tgctgattct 120
ttttccgtaa cgatacggaa ctttacgaat tacgtaacga tacttggttt ctattcgtaa 180
tgacacgaca ccttacggat tacgtaatca tcccttcctt atcttacgaa atgttatgat 240

actttacgga ttgcgcataa acactttctt ttgacttccg acatgtcacg aaacttcacg 300
gattgtcaa cagtgcattc tttagacttc cagcatgtca cggaactgca cagattgcct 360
aacgatgtgt gctaactacc tacgagggt catacgaggg tctcatccca ct 412

<210> 1077
<211> 382
<212> DNA
<213> Glycine max

<400> 1077

acacagacca ataccacaac tttccttact caaatacccc agtaacattt tttcgttcc 60
aatttggca ccgttggatc gactcgaaaa ttttactgga ggtccctagt acataagtct 120
acatttgac cggtggatc tgctacaaaa cgtccataac ccaatatgtt caacccttc 180
cacaaccaggc aatgcataag catttctgc acaaggcaca aattatgctg cacattcaa 240
cagcaaaattt ctgcataata gtgcagattt tcgaaatcac tcttgcccttc ttccaatgtt 300
gcccaaattt gaccctacaa gtcctatatc aagtataat catacctaaa ccacagacaa 360
gcttcagacc aaagcaattt aa 382

<210> 1078
<211> 434
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1078

agtttgtcca ctacttgaaa tgacttagtn taatttataa ataacatgcc attgcagata 60
agatatactt atattcagtt ctgttagatgt tggcgct gatctcccat tagaattaga 120
gttctaccct gataatccat cttagtcctt taatgcagac tatattttgt agattaccaa 180
aaatgaaact ctgattctaa ttgaataaca acatcaacag ccaacaatac caaacaaacc 240
ttatctaagt ttatattcttgg aaaatatcac cctgtatcac aagttcacag ctatgatatt 300
gccataattt ataaaactgac ctgggataag aatggctgg atattatctc atgcgatgtt 360
aatgaaccag acaagtttattt attccttact ccaccaagct aagaacggcc cactagttt 420
attttcaaca catg 434

<210> 1079
<211> 440
<212> DNA
<213> Glycine max

<400> 1079

agcttctttg agaagctaga tccttatcta tccataccccc tctattaact aaattaattt 60
ccttaaaaat aattacggat gaaaataacg caacaataaa tcaaacatca aacataatta 120
ctaataatat atagatatat atatcagggt gttacatcgac cacctgcaca acctaaggcg 180
cccgccccca tccagaggga ggctcccaa gctccggctc caaccacgac tcgttctgtc 240
ggcaatgcct actttggatc cggttccaat gccatgagga actttccccc gaagccaact 300
ccagaattca ccccaactccc aatgacgtac aaggacctct ttccgtccct catcgccaac 360
caaatggtcg taataactcc cgaaaagatc taccaacccc ctttcccaa gtggtatgat 420
cttaatgcaa ctgcattgt 440

<210> 1080
<211> 432
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1080

agctntgata ttatgataat tattgctaga agatctcatg ccatgatttg gaattcctat 60
tgactccagt ggagtacaat gggggggca acattcatca atataacgtg gaacaagtgc 120
acccagtc当地 gttatagag gactgaggag accatacgaa catgtgcct taggagattg 180
aggagtactg agagcttcag aggcaattgg agaagacgtc tgagaaggag agtgagattc 240
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cctagtttc ttcatgttatt ttgatgttatt ctaagagagg tttctccaca tatatgttatt 360
ttgttgaagc aaagatgtaa aaattgattc tgcttacttt catattatca tgggtgcgt 420
tcgtattatt tc 432

<210> 1081
<211> 436
<212> DNA
<213> Glycine max

<223> unsure at all n locations
 <400> 1081

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taagctcctt caactgcaca atgctctaa tatttgaaga gtgtccttgt ggaaccttca 60
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ggggtaagta aattatcttc ccatcagacc ttggatgcaa atgtgattgt atacccatat 180
cagctaaatc ttgacaggta ttcaagccat ccttcatactt gccttgaatg ttcatgagcg 240
ttccaaccac actgatacaa acatTTTCT ccacatgcat aacatcaata caatgtctaa 300
cgtcaagatt acacctgttc ggaagatcaa agataatgga cctttcttg catatgcaac 360
tctgactttt atccttcttt tgagtacttc caaatgcagt atntaagtga tgaacccgct 420
gatataacctg ctcacc 436
  
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<210> 1082
 <211> 456
 <212> DNA
 <213> Glycine max

<400> 1082

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tctgagtcac ctgccgcacg caagctgaac ccaaccgatg agttgtgaac ttttactgtg 60
agtgaacgac tagctgtgac taataatctt tgcatcaatt tgtgaatttt agaatgaaat 120
ttataaatga ggacttgatg aaggccatga ttgtgcatac acaagcctt tgaccaaaaa 180
gcttaccttgc aattataatt gtatcatttgc cacccttgc gagctgaatg attttgc 240
taattgaacc ctaaacctga atgattatct ccagatacct tgtagtgcattt ctaggagac 300
atatggttca aggcattacc ccaaattttgc gggagtggaa ctaattggaa tgcaaaagaaa 360
gagataaaagc atcagcacac aacaaataag ttgtgtgtt aaaaagaaaaa aagacaaaga 420
aagcaatcga aggaaatgtg tggtgatgtataagg 456
  
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<210> 1083
 <211> 432
 <212> DNA
 <213> Glycine max

<400> 1083

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agcttatgag catgagggtt agcctttct tgactataaa aataacgttgc tttagttccc 60
  
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ttgcttaactc ccataataacc aggagcccttg tgcataatcg tttttataaca taagctaattg 120
agtttctgtt gtaacaaaag tgtttgcact acactatgg tcactacatt cgcggtacgg 180
gctaaggctg attcatttat gcaggttgc attggcaag ctgctgtgca caggtgtaat 240
gcaactgtca acttccttga tgaaacaaga ccttcataatc ctgcaactgt aaataatgg 300
gacttgcaca agctatctgt tgatgttagct ggcaatttgc ttggcaccaa taatgttaat 360
attgacaaga cacctatcat ggccgctgaa gactatgcat tctatcaaga ggtcataacct 420
ggctacttca tc 432

<210> 1084
<211> 433
<212> DNA
<213> Glycine max

<400> 1084

agttccggaa gatagtgtatg aggtacattc cctataggca gagttgaaa gagcctgcgt 60
agtcgaagag aagttcaagt ccatagccat caaagtctga aaagagtatg atgaactaag 120
ggatgtcaat atggccaccg atgaaggcctt ggaatgagaa accaagaagg cccgaaagga 180
agaacacgac caaagcaaag ttttggggg ctttataggg cagcaatagt gagctcaaac 240
tccgaagagg tgaaaagaat catcacgggt caaaggcatg atctggaaagg acgagctaaa 300
agcttgcctt aggtcgaaaa gaaatttgc ccaacagtta aagttagact gaaggaaata 360
tgtggccat catcgatgag tgcaaagaga agctaaatct agcggcaact cacgagcaaa 420
ggctatagga tga 433

<210> 1085
<211> 428
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1085

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ctttcatctc ttttcctttt agaatccctt tcatgattat gtatctttc taagaaatct 120
gaaacatcat ctaaacacatt ctttctgga gaaataacat tagactcatc aaaggaaaca 180
tgaatggatt cttcaatatt catagttctc ttattgtata ttctatatgc ttactatgc 240

aaggaataac caaggaagat tccttcattt gccttggcat caaactttcc taagtttct 300
tttccattat ttaatacataaa acatttgcaa ccaaagacat gtaaatgtga aatgtttgt 360
tttctaccat tgaataattc gtaaggagt ttctttaaga tgggtcttat taaagccta 420
tttaagat 428

<210> 1086
<211> 442
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1086

agcttgcaaa tggcattata cattaccatg tgactattt ttttgttata tatcagtata 60
ccatgtgacc acgagttcaa actcaattt a gccaatcac ggtagaagat tttatggta 120
aaaatagtag aagaattaat ggcaagatga ttgattggtg taactacaat aaacagtccc 180
cccactttgg cttaatctg ttntttttt tttttcaat cttccactc caaagttagtc 240
atgcaatatg ccaccaaata ggagaaagtc tgcaaatctc cttaattact ggttggcaag 300
tggagttccat ttgacccacc ttacctgagc aatggccaca gctacttgtc cttttcaagt 360
tctggtaaa agggtgactc ttcatgatca tatcccattt gatatcatat tggcctaaaa 420
gcataaaatg acataaacatc ta 442

<210> 1087
<211> 455
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1087

cgaaccanaa ccggtgaaag tgtgattnta aactgtgagt gaatgactag ctgtgagtaa 60
taatcttgc atcaatctct gaattttaga atgaaatgtt taaatgagga catgattaag 120
gctatgattt gttatataca agccatttga ccaaaaaact tacctgaaa ttataattgt 180
atcatttgca ccctttgtga gctgaatgaa tttgtcaata attgaaccct gaacctaaat 240
gattatctct agataccatg ttttagattct aggagagcat atggttcaag gaaaatttac 300
cccatatttgc ggggagtggaa actgatttggg atgcaaagaa aaagataaag cattagcaca 360

<212> DNA
<213> Glycine max

<400> 1090

tgagctctga tggcgccag cccaccatct tttcatagta gagtaccgat tatgtgtcta 60
ccatcacgat ttcgtctcc ctttccatta ttgggggtac cacctgagcc gccagatccc 120
tccacctttt gggcgtgttc tttgaatgat ccgtccccct tttgcacat gttctgttagt 180
tgcacccat ccggaaccat atcacaattt tactgatact gcctaacaaa ggcaaccatt 240
aggtccttcc aagaatggac tcggaaagg tccaagttag tgtaccaggt aacagctacc 300
ccagtaagac tttcttgaa ggaatgtatt agcaattcct catctttgc gtattcccc 360
atcttctgac aatacatctt tagatggttc ttggacaag tagtccccct gtacttgtca 420
aagtccagca ctttgaactt gggaaatgacc atgtttgggt atta 464

<210> 1091
<211> 449
<212> DNA
<213> Glycine max

<400> 1091

ctgcagcttgc caagtttat aaagaaaactc ttaaatctct cttttatctt tagattttaa 60
catgttctta gttacattca attgtgcattc atcgattcgt tttgttattc ctagcgaaca 120
ttttgttta tattgtatgt tcacaactgt gtaggtatt gtcgttgata aagttttaac 180
cttcattttt cacatccccct atatttcttag tcttacaaaa tgtcatttct cttgtatcta 240
ttaatttttcaaacttttta ttaaaaattt attatgagcg tgtttcgatc cgccaccaag 300
gcccttggac gtgcgtattt catggtaaa catgagaaag tcagttgacc gtgtatttt 360
gcttctccac tatactcgat tcacccgtc gaaatcgatt ttgaagcaca gcatgggtga 420
ggcagcttcc acgtcaagtt aaaattgtt 449

<210> 1092
<211> 436
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1092

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cttttaattg aaagtttggc acgatgctta gtgttgaatt ctggcaatgc taaggtggc 120
ctctgaacat ggacaaatata tctgttaatg gaaattctgg ctccttcaag ggtgggtgg 180
gggttctgat actgcctttt gagtagtggg ttttgttgc gcccctttaga gtaaactctc 240
acaatttcac aatccgattc tacaaataaa gtttatggaa cttccacgat tctacgtaaa 300
atcgagagtt taacaaccat gattggaaatg atcccgctc tctatctttt tatatacccc 360
attaataaaag ataataattt ttgatcagca gactgcctgc gttaagtgtt attactgctg 420
gaaatgcaga tattat 436

<210> 1093
<211> 425
<212> DNA
<213> Glycine max

<400> 1093
atttatttaa acctcccaact gggcatthaag ttggcctac ttggttaaaa ggaaaaatata 60
atatttttc atagaatggg gaacaactaa ctcaaattaa tttacaatga atatttaaaa 120
atgtggagtg actgagttaa aatggattca cttaatcatg ctatatcgag tgactgagtt 180
aaaaatttag agctatatat gtcgctagac gcttatcaac agtataattt tctggtttat 240
aaattaaagt aaacgacaag aaattacttc ttatacattt tgcctacgtg caatcacaac 300
gatacaaaat tccatattat aagaattgcg agtataacta tgttttcatt gagcatttat 360
ttattaaatgg tattgtgcta cgcatgtga ggctgtgcga caagttcatc agatataaaat 420
attgt 425

<210> 1094
<211> 192
<212> DNA
<213> Glycine max

<400> 1094
tgaagctcca tagctacgat tgacgccaat gactgtaact aacatgacta ccatactgg 60
caggactgcc tccggatggc attgaggctt tctaatacat acaccagatt gaactcatgc 120
aagctgttca cttgtcgcc atgttagaggc tgctgttagac gcagtgaacg atgattctt 180

actaaatgat ct

192

<210> 1095
<211> 429
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1095

agcttganga ttatgggta cccgtcatat gtggatttag gtggcgatcg ggcgatggtg 60
aaaatcaact atccccacatc cacgaatcaa acatgaactc accatcccc gttgccacc 120
ttcaactaag ctcacgtact cctacgtacg ccttatacctc gttcctttca acaccgggtc 180
ctcatcaacc cctccaagct tccacaatat ccaagcgatt caatttccaa atatcatgaa 240
ctatcctaaa ccaagaaaaac agggcagagg cagaaaaactc tgcccaaaac acattcacat 300
attacaactg tccttactca aagaccccag taacattctc ttctgtccgg tacgttaacc 360
ataggatcaa attgaaagtt ttactggagg ttccctagtagtac ataaatctac atttgtgaccg 420
ttggatct 429

<210> 1096
<211> 334
<212> DNA
<213> Glycine max

<400> 1096

agagatctac aaacattgtt gtgcctacaa cactactact aacgaggaat gtggtagtgt 60
attacaccga gttttagttag tactaggagg aactcgagag aagactctgg gatgagaagt 120
taactgattt tgcagacgtat cgcatagaca ttgttattat gaaggaatct tacgccaacc 180
tctatgaccc caaggtaaa tcacttattc tggtaaggt gagaggacat ctaacgaagt 240
ttgatgaaga ctgcttgaac acattttagg agaccccgat gactatggaa gagggggaga 300
atttgtgtgc tgattccaag tttgcactcc tgag 334

<210> 1097
<211> 439
<212> DNA
<213> Glycine max

<223> unsure at all n locations

<400> 1097

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ttaacctagg gaattaaaac aaactaaatg gctgagtgt aactgaaattg ttggcaacca 120
aaagtcaccc ccaacagcca acaagtcagc caccattgg tctccaaaa ggctgatgcc 180
taggttgcca attgggcct tattacaact tgaactaaag cccttttagt tgattaaccc 240
aaaacatatt tttggtcagc caactttaca aggattggc cattatttag acaaactaaa 300
cactctaaaa ttgaaataaa gtggtgtcat ttagtcctcc atttggcca tgataacaact 360
cacaacctt gactttctc cttgaaactt gggcttgtat tcaaatacgta tggacagcac 420
ttgttgaaga gcgtccttg 439

<210> 1098

<211> 380

<212> DNA

<213> Glycine max

<400> 1098

aggctctgag caaattcaaa cgacaataac ttttactca aatgaccgct tgagtcccgt 60
agtacatcga gatgctcgta atagaaaagg gaagctctga gaaaatgaaa cgaccattac 120
ttataactac gatgtcggat agagccccgg aaaatattga gacgctcaac attgaaaaca 180
gaagctctta ggatattcca acgacaatat agtttactc ggatgttcga ctgtgtcccg 240
taatataatcg agactctcgc aaatgacaag agaagctctg cgaaaattcg aacgacaata 300
acttctgact ctatgttccg cttgtgtccc gtaatataatc gagaggctcg ttatagaaaa 360
ggaaagctct atgataaact 380

<210> 1099

<211> 149

<212> DNA

<213> Glycine max

<400> 1099

catgcgtgct gggattgatc tgatgcctgc cttactaacg ccatagacgt actcttatcg 60
acatggctcc ccatcgact caagtgcata acatggaaa taaagagaga gatagctcta 120
tacactacgg actatggcgt agagacctg 149

<210> 1100
<211> 376
<212> DNA
<213> Glycine max

<400> 1100

catgagctag tgaacactct cttcgacatc atgtctgtgc tcataatagt gatcatgacg 60
tgcattgaga tggccctgct caactcgcac gtcgtgcact gtgataacctc gctcctgaag 120
acgcttcttt ctgatcttcg atggcactac actcttgagg ggaacattct gaagaactgc 180
ctaccttgtc ttcatgttcc ctctgacgct ggtagcgtc tttggagagg ttggacatac 240
ctctcctgaa gatatgatac gcattgtacc tcacttgtag acagggAAC aagtacacaga 300
caccctcca tgctagccca gagttggtcc caatttagcct ttcattttct acgcacgagc 360
agtgacccctg tggcg 376

<210> 1101
<211> 438
<212> DNA
<213> Glycine max

<400> 1101

agcttatcga gaaaagaaaat tgtataatgt ttgtttacaa cattgttaag ttcaactaaa 60
accctttgtt gaggcattatt cccaaatgtct gtaagaccaa ctgtaaaaga aaaaaaatta 120
aacacttgac aatggatgca tgcactacta tcactatacc agctagcttc attcgctct 180
ttcaaggcatc tatagcaatt ctttgcataa aaatcttgaa actaacactt ggacagctag 240
atctaaccgt tggtgtcgga gtgtgaccaa attaatggtt atatattatta tgaataattg 300
aatattaaaa tactcttggc agtgcataacc tacaaagctc acttggac aaaaacatta 360
cggtcttaa tggatagata agaattaaaa tcaatcaaag taaagatcag ggaggatcat 420
catcaatttt cagcaccc 438

<210> 1102
<211> 447
<212> DNA
<213> Glycine max

<400> 1102

tacggaccta tgatactcg cttgcttcca caaatagtct cggccgaaag acgctgacat 60
cttctggaaa ggtgcagatg accacattgg tctctgcgtg tcatcgact tggggctcc 120
gaataacgag gtgcggataa ccgtaaagtg ctctgcgtc catcgaactc ttgggtcgat 180
ggatagcaag aaggtgacac taaatagtct cagtcgaaag acgctcacag ctccaggaag 240
agtgcagatt accacattgg tctctacgtg tcattggact tgggggtcctt gaatgtatgag 300
gtgctaataa ccgtaaagggtg tctccgcatt ccaccggact cttggccgc tggatagcaa 360
aattgtgaga caaaaattgt ctcgaccgga agatgctgac atctctgtta aaggtgcaga 420
tgaccacatt ggtctccatg tttcatc 447

<210> 1103
<211> 436
<212> DNA
<213> Glycine max

<400> 1103

agtttccttg agaagaactt aactacacac acttttctaa taactaagct caccccttgg 60
agaagcttcc ttaagaaggt tccttagagaa gctagagctt cactacacat accccctctaa 120
tagttgagct caccccttgg agatgagaag ttagagctt actacataca ccccccataaa 180
tagctaagct cacccctcatg ccaaaataca tggaaaaaca aaaaagtccc tactacaaag 240
actagtcaaa attccctgaa atacaaggct aaaaccctat actaatcaaa tggccaaaat 300
gcaaggcaca caagaaggaa aaacctattc taatatttat aaagaagagt ggacccaacc 360
ttggcccatg ggctcagaga tctaccctga gggtcatgtat aacgcttaggg ccttctgttg 420
cagctctagc ccaatc 436

<210> 1104
<211> 429
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1104

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agtttcatga atcaagattc aagagcaatc aagatcaaga ttcaggaatc aagaaaagac 120
tcaatcaaga taagtactaa aagttttcg taacattgag tggcacaaga attttcaca 180

aaatctttaa cgagagagtt ctacttctg gtaatcgatt accgagagcc aacattggtt 240
ttcaaaaactg atttacaaag cttgtaatcg attaccatga gcatgtaatc gattacaat 300
attgtaaaat gttagatttc aaatctcaag agtcacaact agtgataaaa cattgtcaaa 360
tcattgtaaa cttgtcta atgattacaca atacttgtaa tcgattacca gagtttctaa 420
acggtttga 429

<210> 1105
<211> 430
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1105

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ggtgattttc caccatggag atggagcaga agacaaagga gaagaggtga gagaaggcac 120
tatccactag ggaataagcc atggaagaag gagttcacc accaagatga gcttaggata 180
agaagcttgg agaggatgct tcaatggagg aaaagaaaaga aagagagaaa gaaagagggg 240
gagcacgaaa ttgaaggaag aaaaagggag agaagttgaa ctttgagttg tgtctcacaa 300
gactctcatt tatcaaagtt acaacaagtgc ttacacatgc ttcttattt agacttaggta 360
gcttccttga gaagcttagag cttagctaca cacaccctc tcataactaa gtcacacctcc 420
ttgagaagct 430

<210> 1106
<211> 429
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1106

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actctcagcc acttatgata gccggccatg atcccattac ggcttccccct aagctctctg 120
tcctttcttc acgccgcatac ccatgccttg cgaactcctt ggagtaccct cgcgttgtgg 180
tcactgaaac cccgtgcgtat gaaaggcgtg atgcttcgt ctgatggcac tcctctcatg 240
aggttagccaa gctgtcttat ggcgaggacg ggattataat taatacaacc cttgttccc 300

atcaagggaa catttggaca tccttcgcata gaagatagaa tcctgattct tccttcctc 360
tagcgaggga accaattaac agacccccct ccatgctagc caagagttgg tcccaattcg 420
ccttcctt 429

<210> 1107
<211> 438
<212> DNA
<213> Glycine max

<400> 1107

agcttgcct tggttaaac atgattggta catgatttg gacttgtatg tattaatttg 60
ggaaaaattt gatggggaa agactggtt tcgaaatctg cacttatgc agaattttgc 120
tgttgaaatg tgcagcagaa tttgtataa gtgcagaaaa atgcttgtt atggctggtt 180
gtaaaaagg tagtacatat gggttctgg acatttgcta gcagatccca acggtaaaaa 240
tttacaccta tgtactagag acttccggta aaattttaga gtcgatccga cggttaacga 300
attggaacga agaaaatgtt actagagtat ttgtatgtga aaagctgtga ttttgagttg 360
tgcttggc agagtgtctg ctttgcct gttctgctt gttgtttag tacatgatga 420
tggatgtgg aattacct 438

<210> 1108
<211> 469
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1108

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gtagagcaat tatgacctct ccagcaacag atacaacctt ggatggagga atcaccctaa 120
cctcagatgg tccagccctc agcaacaaca gcagcagcct gtccttcct tccaaaatgc 180
tactggcca agcagaccat acattccttc accaatctca caacagcaac aacctcagaa 240
acaaccaaca gttgaggccc ctccataacc ttccctcgaa gaacttgtga ggcaaattgac 300
tatgcagaac atgcagttc agcaagagac cagagcctcc attcagagct taaccaatca 360
gatggacaa tnggctaccc aattgaatca acaacagtcc cagaattctg aatagctggc 420

cttctcaagc tgtccaaaat cccaaatatg tcagtgccat atcatttag

469

<210> 1109

<211> 585

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 1109

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actcaccatc cgacgctgac acattganga ccgtcgact gtagccgcgc actctataca 120

tgctactgcg ngcatgctag catgcttgag gtgctgttat ggacgctgta tttacaagct 180

tcaatgaggt ccttaatgg cgaatctcca ccacggagat ggagcacaag acaaaggaca 240

agaggcgaga gaacgctcta tgcactaggg aataagccat gtgaagaagg agcttcacca 300

ctcagacgag cctacgataa taatctcga caggatgcta caatggagga caacaaagaa 360

cгатагатаг aaagaggggg agcaccaaат tgtaagaaga aaaagggaga gaagttgaac 420

ttcgacgtgc gcctcacatg actctcattt ataaaagtac cactgaggct acacacgcta 480

ttatatatat actacgaacg cttcttgaga ggcatalogacc taagatacac atcacgcttc 540

ttatagcgaa ggcacccctcc ctcgagaagc tcccttaaca cgacg 585

<210> 1110

<211> 368

<212> DNA

<213> Glycine max

<400> 1110

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catgccttgc gaactccttg gagtaccctc gcgttgtggt cactgaaacc ccgtgcgtatg 120

aaaggcgtga tgcttcgtc tcatggact cctctcatga ggtagccaag ctgtcttatg 180

gcgaggacgg gattataatt aatacaaccc cttgttccca tcaaggaaac atttggacat 240

ccttcgcacatg aagatagaat cctgaatctt ctttcttcta gcgagggacc cattaacaga 300

cggccctcct gctagccaga gttggtccca tttcgccctt cttttcgacc acacgggtgac 360

cttgagcg 368

<210> 1111
<211> 459
<212> DNA
<213> Glycine max

<400> 1111

ctgattgtat ttttagagtga gtatttatgt caactacagt cttgctaggg tcaccacaac 60
caatatataca acctaaaggta tatgagcaca gtcatggtag attttgcacg gacatgttat 120
tcacttgtca aaatttagtcc taattgaaga tagtaaatag gaaaagttgt aaggactacg 180
aatgtggtaa tcagcgctgt caaaaattatg gagcatgtga agttaatgtc ttgagaatga 240
ttacttttc ttaagactga gagaagctag atcagtttt tgaatttgac ttctcttatta 300
tccctttca ttctgtccca tgaaaataac attgcatgac atgacaatttatta tattaatatt 360
taattataac atttaattct aaaaatcatc aaaactgtat aattttaaaa ggacaacaca 420
aggacaagga caaggacagt acaaggaggg acaatagac 459

<210> 1112
<211> 185
<212> DNA
<213> Glycine max

<400> 1112

acaaaagagg atctttgca tatctatctt atgtaccgga agctccacat tcattatgac 60
tgaattatcg ctgatcacat gttggctacc cccatgatcg tgtacctaattt atgtgcctta 120
gccatattta acactacgat aactgactac tacataattt ataccataaa aacctttcca 180
gctgt 185

<210> 1113
<211> 612
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1113

tctctctcta actaccactc aacactcgac acataattct agtttatatt gctataacag 60
atacctgtgt atcgcatact cgcaactctaa ctgcgcaact atctcacatc cctacaccat 120
taacacatctt aatcactcta tatctgttcc acctactaca ctcgataaca catatttctt 180

cttattaaac ctatacatct acaatctcta tgnaactgcg aatcacccta acctcagatg 240
gtccagccct taataacaac atcatcagtc ctgctcactt gctttccaaa catgctacct 300
aggcccaaca caagactcat tacattcctt cctaccaatc acaacaacta gcaactaact 360
ctctagaaac aacacaacac attgaggccc ctctataacc ttccttcgaa gaacttgtga 420
ggcaaatgac tatgcacaac atgcagtttc atcacgatac caatacctcc attcatagct 480
taaacaancc gatgggacaa ttggctaccc aattgaatca acaacagtcc cagaattctg 540
aatagctgtt cttcttaagc tcgccaaaat ccataaatgt catgccatat cattgacgtc 600
gctacacaat ct 612

<210> 1114
<211> 394
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1114

tgtgttaagtt attatcattn gaatttctca cgagcttccg ttgttcaatt tcgagcctct 60
cgacatatta tgcgccccgaa tcggacatac gtgtaaaaag ntatgaatat ttgaatttct 120
cgagagttc cgatgtttaa tttcttagcgt attgatattt ataagcttga atcggatata 180
cgtgtgataa gctatgacca tttgaattta tcagagctt ggtgttcaat tctaattctct 240
ctcatataag cgcccgatcg cgcattcgtg tgaatgttat accattgaa ttacttagag 300
ttacgatgct taattcagcg cttgtttata ttcccttgat atccttcttg tataagttga 360
cctctctgcc tgcacatctc tgtgatataa tagg 394

<210> 1115
<211> 372
<212> DNA
<213> Glycine max
<400> 1115

aaagctctcg agaaattcaa atggtcataa ctgttcacac tgacgtccga ttcaaggctta 60
taatatattg atatgctcaa aaataaacat cggaagttct cgagatattc aaatggtcat 120
aatcttcac atggatgtcc gattcggcgc cataatatgt cgagaggctc aaaattgaac 180
aacggaaggt cttgagaaat tcaaattggcataaactttc acacgagatg tcgattaacg 240

cttataatat atcgatacgc tcgaaattga acaacggaac tttccaaaa tatagatggt 300
cataacaatt accatgatag accattctgg gcctctattt tcatcaggta gaataaaca 360
cctagcttt cc 372

<210> 1116
<211> 438
<212> DNA
<213> Glycine max

<400> 1116

agtttatata ccaccagcat cttgtata gggctgttga tggAACCTCT ccaaATGCAA 60
gcttcgcga tgacttacgg aaagatctta gagttgacct tagcagaggt atccatagaa 120
accattgcat cactcaccca atactacgac cagcctttga gatgcttcac attcggagac 180
ttgcaattag taccaccat tgaagaattt gaggaacttc taggatgtcc tctcgaaaa 240
agaaagccat atcttcatc cgggtgtctc cccttttga gcataattgc aactgtggtc 300
taggatacaa caagaggaaa ggaccgcata aaacagactc ggaacggcat agcgggccta 360
ccacggatgt acctagaaga caaggcgagg ggtatggcca atcaaggaga ttgggtcccg 420
tctatggata gttagct 438

<210> 1117
<211> 368
<212> DNA
<213> Glycine max

<400> 1117

atatgcgcac acttccttac atacgttgt tagcacaaga cattatatta accgtaaaat 60
ataatcccc catctacgat caaggcagct ccgacaccta aattattttac acgtacttgc 120
aacgtgtaga tgatacttac atcgacacaca tgtccttggc taaattgaca tacaaggata 180
ctcaaaacat tttgggtac gcaaaattgc acactgtgca cattatggca tttcttaaac 240
ctagacatac actaactcaa tgatgaatct tgactatcta cacaatacgg tgctacatgt 300
catgctctt tcacatttg gctccctaacc accgcatgca aattcaagta tatcatcctt 360
tgctgact 368

| | |
|-------|-------------|
| <210> | 1118 |
| <211> | 494 |
| <212> | DNA |
| <213> | Glycine max |

<223> unsure at all n locations
<400> 1118

cgcgctgacc ttgtgaccct gtgtgcgcga gcttaagaca acccagcatg ctgcttggttt 60
gataaagaac atactacatt gtcgttgagc tgaagagacc cgcattacgt aagcctgtgt 120
cttatatgac acacatatta gcttgactta tatatggcaa gggttggcgc atgaacatata 180
atatgaacaa tggatatgac tgctacgtac gcctctctat ttcatggact tgacatcata 240
cgagttcttc tctccccctt cttgcgtcta agatagctaa cgaatatcac tgacttatgc 300
actgcgacaa gaactgccat acactactga actatcacaa ctctgagtcc aacatataacc 360
ttggttaaca ttattatgaa cttctcaaggc cagggagaac cttgagcaag ataccgagta 420
ccgggatgac aatgacaagg atctgagatc catcttact gggtctatac tgtcatgatc 480
actacgcacg ttan 494

| | |
|-------|-------------|
| <210> | 1119 |
| <211> | 222 |
| <212> | DNA |
| <213> | Glycine max |

<400> 1119

tctcaatatac tgttcttgag tcttaacgc gctctgaccg gttattgaag ccgtgcttgt 60
cgccctgaaag agtgataaaa gaatacccac cgagcatatg tggtgcacac tcatttactc 120
actttacaaa cgaactctgg ccgatcgtgc acgctataac ctacgctaaa ccgctgagag 180
gaaagtatgc aatgttatctt gtaacatqqt tqtttaactqa tc 222

| | |
|-------|-------------|
| <210> | 1120 |
| <211> | 423 |
| <212> | DNA |
| <213> | Glycine max |

<223> unsure at all n locations
<400> 1120

gtatgatacc tgttaacttt atgcccaccc cggatttcga cattctcaac ttgaaacagg 60
gtttaggggt ggggaacat aacaaaagcg tttaatcqqa atqcattgat agttaaga 120

ttttataatg aaatgcaata tatactaata tagtcgcctt tcctagcgat tcttctaata 180
catatatttg agatgattat gtaaaaatca ttatattaaa ttagtaatgt atcaaaaacta 240
aaattctaaa tatatgttga ggcatgactt aatttatgtt attttatcaa aataaaactct 300
aaaatttatt ttaagaagct ttaaggtaa cactataata taaactatnt agtgatacta 360
aactcgctca ttcatgatta tttgtcgcgt tacgaattca cttttactat taactcaaaa 420
agt 423

<210> 1121
<211> 412
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1121

ttcatggct aagtaatata aaaagctctc tctgatatacg taccagattt cttacattgt 60
aaagcttcat acgatacata cttanaggtac ccaactttgg ccataatggc caatattact 120
aaacaataga aaggattta taaagttatt tctcagaact gaatagatta gagctaaacg 180
actatctatt acctgagata tggtgatgac aattacgtct acacaaattt tgaatatacg 240
ttcttcatt gttgaaatca attaaaaattt ataaaaattt gtcaatatta ttgcttccta 300
attaatactt tctataattc tataattgtat aatatgtcg aactaactaa taaaacagaa 360
tgtttataa tatactacca tgtagaaat acttatatac tatcagtttc at 412

<210> 1122
<211> 392
<212> DNA
<213> Glycine max
<400> 1122

agctagacca atcctgaccc aaccggca tagttaatca gtgataacct gtgatgtacc 60
taaacaggcg agtcctggc agtcaaccga taagggAAC aaagaccaca aagcatggag 120
gcttgtgtgg aggttggcca gctgcggaaac tttgattgtat atatggata tggactctgg 180
taatcgatta ccaagggcg taatcgatta caaggctaa aaagtaagac aggagactaa 240
gatggctct ggtaatcgat taccaggctt gaaaactaga tcatgaagct tggaggcctt 300

ctcgtaatcg attaccatgg cgtgtaatcg actaccaggc ttataaatga gacttgaatg 360

ttgaaggagc ctctggaaat cgataccaag ct 392

<210> 1123

<211> 379

<212> DNA

<213> Glycine max

<400> 1123

tggtaggta gccatggaaa agcagagcgt ttggatgtat ttcgtaaatt tcagaaggct 60

attgtgaaat gctggtaaaa acacgaatgc caagcagata taaatttcaa tgaggaatgt 120

atagggtcgt gtgaagcaac ggtcgaattt tccttggttc agtagtgaac gtgctattaa 180

tgttaagtga ttcgtttggg cacgttcaga ttgctgttagt tgctataatt cctctagcac 240

acaaatgccc agcttgcccc tcagttgttc aaactgattt gcatccaaag cctttgtgaa 300

aatatctgct attctgtcct caatgtcaac atgcttcagc gtgatcaattt tatcatcaac 360

aagatctctg atataagtgg 379

<210> 1124

<211> 449

<212> DNA

<213> Glycine max

<400> 1124

agtttcgtcc gcagatccct catgtaagac tatgtctaaa ctattcaaca ttatgtaa 60

acataattaa aacccaaaact taacccgcag atccctcatg taagactaag ttttgcatt 120

gcttcaatca agttctaagg caacagtaca ttcccataatg ctaaagtcat ctaactgtga 180

acacaaatgg gtgatcagac caaaagcata ctaacatcaa gcattgaagg aagcattgaa 240

cacagaatac acaatcaatt agttttagg tatttacatc atctgttcat ttgaaatccc 300

caacttagggt gttccgccac ccattacaga agagacccta tcaataatta gcttactaac 360

ccttaggtatc tctgcaaaag ctgctcctct tgctacccctt agagctccctt tccctaaata 420

ggcaatgtgg ctgctgtgga attttgc 449

<210> 1125

<211> 459

<212> DNA

<213> Glycine max

<400> 1125

cgccacccagg ctcgcccagg cgagcaaggt ttcttcctcc agaaggcaaca gccttctgga 60
ggaatcttct ggagggccca agtgggtctg gttgctattt gcaccccat tttactaaa 120
tacacccttgcctttt ggtgattctt tttcgtaaa gttacggaaa cttacgaatt 180
tcgtaacgat acttgtttc tttcgtaat gttacggaac cttgcggatt acataatcat 240
ccctttttg acttacggaa tgttacgaaa cctcactaat tgtgcaacga tgcttccttt 300
tgatttccgg tgtgtcacgg aaccttacgg attgtgcac aatattttct tttgatttcc 360
ggcacgtcac ggaatttcac aaattgccta ctgatgggtg ccaaggcacct taaaaatgac 420
caaacacaag ttgcatgccca ccaagcacag gtccctgga 459

<210> 1126

<211> 461

<212> DNA

<213> Glycine max

<400> 1126

gtcacctgca gctgcagcta tcacgcctc acacaatact ttgtatgtat attacaccag 60
agatgtttgt ttcacccagg gtaaaacaat atggagttat caagcatttgc cccggaaattc 120
aaatgatggg tcaggacgag gctgaatgtat tattccatgt cctatggcca attgaacagt 180
cccttgaca agtggcgac acaaggatttca caaggatttca tattttttca ttttccccaa 240
gatacagagt gtccttatatg agagagacag gaattgtatg gttaaggac agaccatcac 300
ccatgcaaca gtcaccaggat catacataga cactattatgt atatttcttctt cttacatcat 360
aacactgccc tattcttaag agataaaaaaa gggaaaggat ccagaatcag gacgcaatac 420
ttcacatttg ttcataccaa ttcatattgg cctctgctca a 461

<210> 1127

<211> 449

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 1127

tatctggtca aggtgaaata aaaatggtct ttcaaaactaa tagttgccag attcaatattt 60

ctatattta atccaagtga acactccat aaggtttgt tcagtttcta tcaagaacat 120
acatcaaata aaggatgcac agctttaga taaaaagttt atgaactcca acataacctgc 180
aaaagacaac tacattcgac tcgtttttt gatacactgc cacaggtctt tataacattc 240
ctgtagaaca gcttctcaac aacaatgaca ttcagatggt tgaaaatatc aaaaccagac 300
tgcttgagtt tatcatatTTT atcaatatga aatttatgaa tgtagcgctg ggcatatggc 360
agagtccagt ttaccaatga ctntttcaaa ctacaatcag caagaccata atatattgg 420
tcccgagttt ccacccgtcac aaagtctata 449

<210> 1128
<211> 457
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1128

ctcatcttta ttccgttcat tctcgatttc ttttcttttgc tcttaacgc gcttttacgg 60
tttatttaag ccgttttctc acctaataaa tgataaaatg aatttcaacc gatcattttgt 120
gttgtaatct cattcaatca ctttaaaac gaaatctaac cgatcggtca cgctataacc 180
tcggtaaac cgaaaaaaagt aaaataatca aaatatcttg aaaaataata ataaaataat 240
caaaatatct tcgaataaaaa taatcaaaaa aatcaatcgg acgttnntct ttggaaagttt 300
ccttgaatga attgattaat aactaaagtt aaattaagac taaaatcaac tcacaaatca 360
agttttgtcc gaaaaatcac taaaaaccgt tttaaggtcc aacgccttaa gcggtcctct 420
ttgcttttat cggttaacat ggaccgttca aaagcat 457

<210> 1129
<211> 338
<212> DNA
<213> Glycine max

<400> 1129

atgacaatgc ttaccaagtgc gagctggccg gtgagtataa tggtaattcc aactacaatg 60
tctctgattc atctctttgt gatgcatatg gagaatccga tatgtatgact aatacttctc 120
atgacggaga gaatgtatgat gacgtgacca caagcaacgg cagggatcca cttAACGACT 180

| | | | | | | |
|-----------------------|---------------------------|-------------|-------------|--------------|-------------|-----|
| tgtatgacct atgacaaggg | ctacagcaag | gaaaggctag | gacgctcttc | tacaattgtc | 240 | |
| gtccatacta | tgcaataca | agctcaagtt | tgaagagaaa | aagtccatgt | tgtgacttga | 300 |
| tcatggcca | tatggaagac | taaataaacac | cactttat | | 338 | |
| <210> | 1130 | | | | | |
| <211> | 414 | | | | | |
| <212> | DNA | | | | | |
| <213> | Glycine max | | | | | |
| <223> | unsure at all n locations | | | | | |
| <400> | 1130 | | | | | |
| ttctacaatc | tcgaggccttc | anataaccct | cctcaccttc | ttccgcacatca | ttatcatcag | 60 |
| tatgaggacg | aatattacat | tcttctctat | gccaatttc | agattgatct | atccggtttt | 120 |
| cctcattggg | ctccacactca | atgctatcct | tttctgagct | agcatggacc | accttatctt | 180 |
| ctgcacgtgc | ctttgtctta | taaaccattt | cagactcatt | aaaaataaca | tcatgactta | 240 |
| taatgcacatct | tttgtgtcct | ggctctaaac | accacaaatct | gtaccctta | aaaccctgag | 300 |
| gatatcctat | aaacatacac | ttgatagctc | taggttccaa | tgtgtcttgc | cttatgtgag | 360 |
| cataagcaac | acatccaaac | accctacgtc | tatcattatt | tggaggatgc | cctg | 414 |
| <210> | 1131 | | | | | |
| <211> | 435 | | | | | |
| <212> | DNA | | | | | |
| <213> | Glycine max | | | | | |
| <223> | unsure at all n locations | | | | | |
| <400> | 1131 | | | | | |
| ctcatctaca | ctgtacttct | atataatttg | tatgacatag | tgcacatgt | ataatgtatgc | 60 |
| ctatagtc | gagatgaggc | atatacaaat | tgctatcttgc | aaaacctaag | accatgcccc | 120 |
| ttaatgtat | aaggggacat | aacaatcaac | atgcaaagga | taaatagtat | acatgacaaa | 180 |
| cctgtactcc | aaatgtcgga | cacttactt | tggcttctac | aaaagccaaag | tncaatttagg | 240 |
| atgtacccaa | aaataaaagac | ctaaattaat | caacagtcaa | acatcaccta | ctaataaaaga | 300 |
| atacaccgac | aatatggtag | ggaaccacac | aaataaaagtt | ggcatttaaa | gtcacccaaag | 360 |
| agatgaagag | aaaacatata | ttgccatcat | ctgctccatc | atatggtaaa | cacactaaag | 420 |
| ttaggaaagg | aatga | | | | | 435 |

<210> 1132
<211> 438
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1132

agctttcttg cttcttagat ntcattcctg ttcttggta ctgttatcg ccattcctaa 60
aactgccaca tcacaactta gatcttaatt gaattttga gcttaccgaa ccatattaaa 120
taatttcaga ttagggacta tactgattat ttcaactgtt gtttcataat ggtgtttatt 180
tgtgcattgtt aatctccctc atgacattct gtattcatac attgtatcaa cgtaacaaag 240
aaaacagtat tattgtgttg tcgaccaatg ccacaattaa agagggccac cctgtactat 300
gatataatgt tacttatgtt gtcttcttt cacaacaaat atgttggat attatttat 360
tatctatgtt gttgtttggat tattcatatt tcttcctggat attgattct tcagctggag 420
acgagttctt cttacata 438

<210> 1133
<211> 380
<212> DNA
<213> Glycine max

<400> 1133

aaccctacat ggagtgattt gctacatcta aatagtaata acatagacgt ctgttatgcc 60
gctttaaaca cttaaatatt ataaacatac tcaaagagaa tgctgctcct tatctataat 120
taagggttca atatgctcct tctacctgct tttaattcac tctgattcat tcaagtgtgc 180
ctactataacc acatgactat attagatacc tgtctgacag cttttctact tccactacct 240
aactatatta attatgtgtc atgcagcatg actcatatac tatcacgtgt cattaagatg 300
acatagtttag tatgcattggc aatattgact tgatctaata cggatcacag tgaatcttc 360
aaaacttaag accactttat 380

<210> 1134
<211> 380
<212> DNA
<213> Glycine max

<400> 1134

acatcttaca tagcgagttg atctctctct tagcgccgg ccataaatct tgcgtcttc 60
cagataactct catgctgcta agcacgctgt atctacgctt aacggtagat gctagctgag 120
cccactggat ccgcttatcg cgactgctcc ttgtgaaagc aatgacttcc aagagtattt 180
tgcgtatgcc aaagaatcaa gagtcaagca cgttccattt aatcatgact ctggttgctg 240
gaagaacctt gtttctgaga ttcacgattc aagaataatc aagtctcaag aggcaatcag 300
gtctcatgaa taatcaagac catgagtc aaactgttga tcaagaccca tgagaagact 360
caatcgggat aagcactaaa 380

<210> 1135
<211> 333
<212> DNA
<213> Glycine max

<400> 1135

tgcctaacag gccaaaccttac aacagcatgt cccaaaggatc tcagctttat gatgcacata 60
ccaaagttga ctatgtgaaa ggatgttatg accaagtgaa ggtgccattt gtcaagaaga 120
atgatggcta taccttagcat gcctacatga tgcgtacga agtggttctt gaacacgatg 180
atgatccctgt acatttgatg gctaatgtgc ttcaagaatg atgaaatgat gagaatactg 240
attctgacca aatacaggct tcaggcgctt gaggagatgg acacacccct ggagtggaga 300
atgatgaatg cccatatgga gaatgataat ggc 333

<210> 1136
<211> 433
<212> DNA
<213> Glycine max

<400> 1136

agcttgtat gttgttgttta gttggtttgg cctctggctc tttcaattat agggacagta 60
ttttgtattt ttttatgttc tgtaatttcc ttggctctgg tacaagtttcaactaatgc 120
tctcatgaat gaaatataact gttatttaggt gatccttgaa cagataacca gttccccatg 180
gaacaacttt ttgttcatga tgcgtatgg cttggttata gaagctatgc tcattcataa 240
aatatgtttt aatttgtatg ttacgttacg cctcaatttac cacataataa ttttggaaata 300
tttccattaa agcttaatgt catctttgt tggtaagtc tcttctgctc tgcacccgac 360

atccctactc aaaacaggaa ccattcattt ttttgctttt ccttaattac ctgctttat 420
gtgtatatca tga 433

<210> 1137
<211> 436
<212> DNA
<213> Glycine max

<400> 1137

attcttaca atcaatctat ctactgaata acaattctaa atgtaagtcc acattcttgt 60
tctttcttg tttgacatgc acatttgctc aacttcatga aagggaaacac aaatctcatc 120
ttaagcatgc attcaattta aaacaaaagtc atacacccgt tttcacaaaa agataaaaagt 180
gtttcactgc catgtcatct aaaataagtt aaactgttca aaatgtttca agataagcat 240
aataattatt catatataaa actagtagta tatatagaca taaagggaaat actgtacgat 300
aacccaaaatt ataataataa taaatcaaaa agtggaaaagt gtcaccagga attaaaattc 360
ctgtgactag tcctgagtct cctgtgttg accatcctcc tcatttgtca gctgaagaac 420
tggagtagtg ggagga 436

<210> 1138
<211> 280
<212> DNA
<213> Glycine max

<400> 1138

gtatgacagt caccgcttta tgagcgcttt acaccagtag cgcttcgagg ccatcaaagg 60
atggtcgttt cgacgggagc gacgcgtcca actcatggac gacgagtata ctgatttcca 120
ggaagagata gggcacccggc ggtggacatc actggttacc cccatggcca agttcgatcc 180
agaaatagtc cttgagtttt atgccaatgc ttggccaaca aaagagggcg tgcgtgacat 240
gatgtcctgg gtaaggggtc agtggatccc gtttgatgcc 280

<210> 1139
<211> 354
<212> DNA
<213> Glycine max

<223> unsure at all n locations

<400> 1139

tctatataag ctgaaccatt ntatcagtgg actcaagtgg tgtttcttc gcaaactaag 60
agtctatctc ttgtatccta ctgagagtga ctctgcataat ttcttgagtg ggtcaagaac 120
accttggctg tcccaaagga ctttatcaac ctttagtgggt cgccctcggt gggacgagag 180
atgctttcct tcggtttatac gtctacccctg ttctttcaaa tcacaattgt agataatacc 240
tcctatgaca ataaatatct tgtggtccta actctccttt tatgcactcc actttcgtga 300
ttgatgggcc tcaaacgaac tcaaaacgat accttgcacc tcataattgga atga 354

<210> 1140

<211> 434

<212> DNA

<213> Glycine max

<400> 1140

agcttaactg ctttgtaaaa cgaaatactc gatctatata atctttgtta tcattaaata 60
tatttcaaac tagttcaatc atatgcatca aagtatgaaa gctttcaaaa aaacatgaaa 120
accttgaagt agtattctaa acaatatttt tctgagtaca atattatgaa aaataacttt 180
ctaagtgtag tagcaacata ataagaatcg ttataacata aactaaattt gtcataataa 240
caatgtttt agagatacat ttatttatgt aatgatcttc taaacaagag caaatgcata 300
ttgacattag gttctcataa tcaagtcaaa cattgaataa tgagtgttat gactaaccac 360
tttagagagct tagttgtctt agtacttgaa cctctatgtc aagaatttct ggacaccaat 420
gtagtcttga ataa 434

<210> 1141

<211> 425

<212> DNA

<213> Glycine max

<400> 1141

atactgcatt gttgactaat tgggtgtgtt gtttattaca tctatttca gactcccaat 60
ttgcagattg agttttgggg ctgctttctt gctgaacaat gtttgggtt gatgtattgt 120
ataatatcct tggcttctat cgccggctgctg tctgttatat ttctttcaaa acttattctg 180
gctctgaatg catacttaag gaaatattat gttatggata atgatataca catctcttta 240

atggaacgat attaccgcgt tttcaacgca tgtgttctca tgtgatgaat tgatgattaa 300
ctgttcgcca atataataaa attattgctc tttactctaa tatatatagg taacgactta 360
atgttgaaga ttagagaagg tgtatccctt ctactccatc accattaact atcagctacc 420
ttctt 425

<210> 1142
<211> 453
<212> DNA
<213> Glycine max

<400> 1142

agctggcttg tggagcttct atggagggtt gatcttcgag cttcaatgag gtcctttaat 60
ggtgatttcc caccatggag atgcagcgga agacaaagga gaagaggtga gaggaggcgc 120
catccactat ggaataagcc atggaagaag gagcttcacc actaagatga gccatggata 180
agaagcttgg agaggatgct tcaatggagg acaagaaaaga gggagagaaa gagagagggg 240
ggagcatgaa attgaaggaa gaaaaaggaa gagaagttga actatgagtt gtgtctcaca 300
agactctcat tcatcaaagt tataacaagt gttcacatg cttctattta tagacttaggt 360
agcttccttg agaagctatc ttgagaaaac ttcccttgaga agcttcttg agaaaaactgt 420
cttgagaagc tagagcttag ctacacacac cct 453

<210> 1143
<211> 459
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1143

catgtcctgg ttacatactt acatagcatt aatcacaaca tacacgccaa taaaatacttg 60
gtaacaagga attataccct gcaaacaag ctatggaaa tggaccacc tcccctaaac 120
ccaagtataa gttaataggg ttaattataa aatggatccc cttatggat tatttcacta 180
aatggatcct ttaagggaaactagaatttgc tctttggag taaaaggat actcttttat 240
cataaaatataa aatcttataat attacattt aagagataat atattatatt ataaaaatata 300
aataaaatataa aaaaatataa ttttgaatta ttttcttaa aacttacatt aatactataat 360
tttgtaacat atattnaat gaacactntg atatgtat aaaagaatataat atttcttatac 420

tcanagatca tgattcatta aaaaataaaag tcatgaaat

459

<210> 1144

<211> 333

<212> DNA

<213> Glycine max

<400> 1144

accacagagt ggtacctgta gatatgtctc gggggcatg agaacctggg gacgtcatgt 60

gggggtgctat tgcccaaaac caagcttgac caatcccgac ccaaccggg catagtcggt 120

cagtgagaac ctgtgatgta cctaaggcagg cgagctcctg caatcaacag ataaacgata 180

acaagaccac aagcatggag gcttgtggtg gctggccagc tgtgaatttt gtgtaatatg 240

tggatggtgg cctctggtaa tcgattacca agggtggta atcgattaca ggcttaaaaa 300

tgaagacagg atgcatagat ggtctctggt aat 333

<210> 1145

<211> 461

<212> DNA

<213> Glycine max

<400> 1145

cggagaagat gttcaatgg aggaaaagaa agagggagag aaagagagag gggggagcac 60

gaaattgaag gaataaaaga ggtatagaag tggactttt aagtatgtct cacaagactc 120

tcattcatca aagttacaac aagtgttaca catgcttcta tttatagact agtagcttc 180

cttgagaagc ttcttgaga aaacttgctt gagaagcttc ttggaaaaa cttccttgag 240

aagctagagc ttagctacac acacccctct cataactaag ctcacccct tgagaagctt 300

ccttaagaag attcctaaag aagctagagc ttagctacac atacccctct aatagctaag 360

ctcacccct tgagatgaga agctagagct tagctacaca cccccataaa tagctaagct 420

caccccccattt acaaacaaca tggaaaataat ataaaagaag t 461

<210> 1146

<211> 347

<212> DNA

<213> Glycine max

<400> 1146

ttaatagtc attgcaccag atctaacctc tgcatcagag gctggaagcc ttctcaaaaa 60
catgaaaacc ttgaagtagc attcctaaca atatttgtga gtgttctata ttgagaaaaaa 120
tgactttcta agttagtag cgacccaatg agaatcgtt taacataaac ggtactcgtc 180
attactacaa agtggtaga gaggcattca tttatgtaat gaagttctag gcaagagcca 240
acgcctattg acattacgtt ctcattatca agtccaaacac tgtatcatga gtgtgatgac 300
taaccactta tagagcttag ggggcttagt acttgaacct ctatgtc 347

<210> 1147
<211> 696
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1147

acgcgtcca cntctcgcc cgtgntgttt antgcanttc attacctccc gtcgacagta 60
atatttgcgtg tgcgttc tctacgctca tgccgtact actcgctcgtc tactccgtc 120
ntagtgcact gactcatant cgtacactct cgctcattnt atcgctatca ctacatgact 180
tactagcgta ctcttatgag tgtctctcac attctctcta ctctcttatac tctatcgat 240
ctttctcatg cacatctctt cttcatctt ccgatacgta ctnnnnccgc ttcctctca 300
cgccatgc acatgtgtgt gctctcgatc atacatatgg ggtagaatat ctcgttagacc 360
tttgtgttgtt cggtagagag gtcagagttg tgccttata caatatatga cttgtttaga 420
acagctgccg tctatgctta aatattatta tagcacatgc tcctgcttcc tcgttgtgt 480
gacgactaca ccaatgtgtgt gacatgctgt atcttgcac acatattcat gtgtactcat 540
gctacgcacg gtccttcac gcgcttcatg ctcatgcata cctaaaatca tcatacacgg 600
tctctcacaa tgtggagtca acccatacac actattatca tacctgcttc tttagaatct 660
attgataacct ttcttggta ccctacagat actcct 696

<210> 1148
<211> 422
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1148

cattatttga ttgaaaatatac tccattataat tttaaaagtc ttactctt tcactgcaat 60
tactcctgct caagtaagat tatctttat gtaaaactca ttcatccatt tggtaaaaca 120
caccatgaca ataagttta ttatcttcaa ttctttaca caatttcatt tgtacaagta 180
tgcatgagtg ttcagattt tagtgcttga gtataaaaaa aaatataatt tttgtcccct 240
tatgtttcta aatctgtaat ttttagtcccc aattntttaa ttgacatatg ttatctctca 300
ctttttaaaaaaa aaaatcataa ttccagatca cccgactaat ttcaaatgtt aattgtgtac 360
tttgtatTTT ttttctttaa attaaactta ttagtaatta aatacataaa aaaaaataact 420
ct 422

<210> 1149
<211> 429
<212> DNA
<213> Glycine max

<400> 1149

tggctgctgt catcatgtgc aagcgtcgca atatattacg ggatTTAGTC agacttccga 60
gtgaaatgtt attgtcgatt gagatagctg cgagcttcgg ttggatatgg cgagcgtctc 120
tatatattgc gggactcaat gagacttact agtggaaatgt gattgtcgta cgcattcgct 180
gctacctatg gaacaacaat tcgagcggct gacatattgc gggactcaga cggacttccg 240
aacgatatga tattgtcgat ataatatgct gagagcctcc gttataaaca tctagcgtat 300
cgatataattt cggcactcag tcagacgtcc tagtgagatg ttaaagtgcgt tcgaagtcgg 360
tacgcgttat ggctattaaat tacgagcgtc acgatattt acgggactca ttcagtcttc 420
cgagtgtatg 429

<210> 1150
<211> 438
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1150

agctcgtagc ctattcgaaac aacaataact nttaactcgg aagtctgatt gagtcctgta 60
atatatcttg acgctcgaat agtaagacccg aagctcgtag cagattcgaac cgacaataac 120

atgtcactcg gaagtcctat tgagtccgt aatatatcga gacgctcgaa ttttaaaacc 180
gaagctcgta gcaaattcga acgacaataa catttcaactc ggaagtccta ttgagtcccg 240
taatatatcg agacgctcg aatTTaaaac cgaagctctg agcatattcg aacgacaata 300
acatttcact cggaagtccg attgagtccc gtaatatatc gagacgctcg aaatttaaaa 360
ccgaagctcg tagctaattc gaacgacaat aacatttcac tcggaagtgt gagtgagtcc 420
cgtaatatat cgagacgc 438

<210> 1151
<211> 508
<212> DNA
<213> Glycine max

<400> 1151

agaacgcgca tgaaccatga gaacatcgag atccgagata ctctccaggc gagctgacgc 60
gtgcgagctg aagatgtcac agtgaacttt ataagcttat gtgctggcca gcatagggg 120
caatcgacta tgtatagcca acgtatttac ttatgagata aggtgtgatg acggtccttc 180
cacattatat tattccatgg tagattgatc gtctccgctc aaaggaacctg cttgttagag 240
tcaagaatat tgagcgtcgg tcaacgttat attataagta gacacctata gaggaactac 300
cacgttataat tgggttaatt gtggtatagg cccctatagt tgatgaatc ggtcattaca 360
ctcatttcct ctagttatgt ctactgaact catgctttat actttacta tgaatggaat 420
ttaacacacc tcttatataa ccgtccaatg tataacaaca ttacgagaca tggatgtcat 480
tatagatgat atgaacctac actgctct 508

<210> 1152
<211> 380
<212> DNA
<213> Glycine max

<400> 1152

aatggatatg gtttaggtgta tgttaatcaa ttagacttta tccgtatcct tggatgtat 60
taccttgaaa actgccatgt agtttgtgaa cagggttcct agtaaggttag ttccaaagac 120
acctttgaa ctgtggacaa ataggatacc tagtataagg cacctgcatt tttgggttg 180
ccaggcagaa ataaggattt ataatccgca agaaagataa ttggatgcaa gaacaatcag 240

tggatatttc attggtttac cagaaaagtt aaaagggtat atgttttatt gttctaatca 300
tagtatgaga attgtcataa ctggaaatgc aaggttcatt ggaaatgatg aaatcagtgg 360
gagtagttt ccacgagaag 380

<210> 1153
<211> 412
<212> DNA
<213> Glycine max

<400> 1153

acactcacac atatacttag ttcaaaattc tatgattaag agcatatatac tactgaaatt 60
aaattcatac acgcacagat gaacacaaca caaaggttc tctgtcagtc ggagttgtaa 120
tttttttaggg tggttataact atgtatagtc atttacaaaa ataccctcca ctgttaaact 180
cttatgactt aattatccct tttatatctc aagatatcta ggatgaccaa taatcaacct 240
taattatccc cacctaattt ctaaccccttac attaatgact aagttttctt tttaaagctt 300
taatttgat atctagacca agatctaattt atttacatct aggtcattaa tgagttgacc 360
attatttgac caagaaaattt ctctaaacta tctttattct atgtagaagc tt 412

<210> 1154
<211> 408
<212> DNA
<213> Glycine max

<400> 1154

tatacaagta ctcttgaggt ttcttccatc agagcctcggtt tattgagctc ttaatcttct 60
ttttcttctt gtattcctta ctggattcgt gcaaacttctt cattcatgga tccaaaatctt 120
cattttcattt cttacaagct tgaaacatca aggatctaag atctttgttc atctaataaa 180
atacatgtat cttcatcaac gtaaagagag tctctccat acttaaaccc taatcttggc 240
gtctttggaa gctaacccttc attgaatgtt gtttagatgt tcaaaatttc atagctactg 300
catatgctgg aactgtatca tgggttgggtt ctcttgtaat cttaacgcaa aaaatgagat 360
atttgagtgc caataacttac gcgttaacctt atatctcacc tacctcat 408

<210> 1155
<211> 392
<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 1155

gcattacatt attactcatg cttctaacca tgtctaatat tggatngattt ctgcgttctg 60
ccacaccatt ttgatctgga gaaccaagca tcttgttattg tgcaaacactc ccatgttctt 120
gaagaaaactt cgcaaatacgaa cctggtgctt gtccatcctc tgtgtatcta ccataggact 180
ccccacccctt atgtgatctc acgatcttaa tatgtttcc acattgtgtc acatctgcat 240
ccttaaaaac tttaaaggca tctaaagctt cattcttaga atgaagtaag taaagacaca 300
tatatcgtga ataatcgaaa ataaaggata tgaataactct ctgacttagtgc gcattcatgt 360
ctgggcaaaag tatgtctgta tgtatgattt ct 392

<210> 1156

<211> 412

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 1156

tgaatgaaac accactgtca ttgagaacaa tggatccatt tttaaaaata atacctaattg 60
tgtattcaca actgttgact gcttgctcat ccagaatgcc atagtttat gaataacttgn 120
gttgatactg aaaccttgc tttcttacag gtttaggttgc tgccatatat atagatgact 180
tttcatatca gtgctgcatt tttaaagat taaaaataca cctgctcatg ctttctgtat 240
gtgttgtcaa ctacaccaat gatgtgacat gctttacctt gcatcaaata tgcattgtgt 300
atcatgctat gcatgaggct ttcacgcgt ttatgttaat gcagacaaaa atatatcata 360
cacggttttc cacaatgtgt atgttactca gaccacaata tatcatacat gc 412

<210> 1157

<211> 425

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 1157

ataaaaaaaaaa tctaattgtat agtcaaaata ctcaaagaaa taactttgtt atgagacatg 60
attagttttt ttttatctat aaaaattaaa gatatatatt aaagaatgtat taaaattcact 120

atggattatt taatcaattg aattaaactc cttaaatata aaatataatc aattaacatt 180
tgacactaac acttaaataa ataaagtaat aactttttt ataaaataca aaagttagtca 240
atctaatcat cgaataattt aatattatct ttgttaattat atataggaag aagtcaaatt 300
acaatcatat gttttcacac caacagttat attgtaaata actctatata aaatataaaat 360
tttattgatt taattgttaa attctaatta tatattatct tgagtgtggg tctcacattc 420
tctan 425

<210> 1158
<211> 436
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1158

tcctggtnaa gtnttatgca aacaactatg cttgacattg ttgtttcta atttgtata 60
tgggatttat attagtttt catattggtg tttcaggat tcctagttag gatcgagat 120
ctcaactctt aaaaggcaga aaaaagtgcgttggaaagctt ttgagacacgc ccacgctatg 180
ggtgttatta tgtttgactt gccaaattgc cctaagaaac gttgtcaact agagacatcc 240
tctgttaatg gagaaggatc atctaccat actgttactg catctttga aactgccctg 300
aggtggcag ataagtagtt gatcgagtca aaactgcatt tataaggctt gcaaattgtc 360
cttcttttag tatcggtgaa ttcgaggaac tactaagaca aattagtaat ctggacttgt 420
ctgagttctc ttca 436

<210> 1159
<211> 426
<212> DNA
<213> Glycine max

<400> 1159

gcacttatgc agtaaagaga ctgtctaatg catgtttaca acatggctaa gttcgactaa 60
aaccctcttg tacactttat gtgcaagtgc tagcagacct actggtgaga aaaacaaatt 120
tcacacttga caatggatgc acgcgcgtgc atcgctatgg ccacataact tcattcgact 180
gttgtaagca tctacagcat gtcttgcaa tataatgtta gatactaaca cttggacagc 240

tgtatctatg cggtgacggc tgagagtgc cacattgatg gatatatcta tcacgaataa 300
gcgttatatta gaagactcga ggccgtgcgt gactacaag tcttagtttg gacagaaaca 360
tgctcggtct gaatggaatg ataagaacta aaatcaatca ctgacatgag cagggatgat 420
catcat 426

<210> 1160
<211> 502
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 1160

cgcgttgca cattgtgacg cggttcgccc gcgcatttg aggcaagtc ggtctacagt 60
catatgcttc tggcactcta cctcatattt gtaatagcca gtgttagttt ataaaggcctc 120
aagctcatga ccgtctatct atgcgcgcgc gtatgatcac ctgcggacg ttgtctggc 180
caccgatgat gtcatggaat gagaagtcat gcttgcctta tctgattacc aacacgcacc 240
cctcctgttg aggtgcttgt gagcgctatc aggtactgat ctcaaacaga gaaagaggaa 300
ttgcaatgtt gcgatcaaa tgcttgagct tcactgccta cctaacagct tgcactgtgt 360
cattcatatg ttctgtccaaac cgtgcctcgta aaccgaacgt aatatgagct ccatcatctg 420
tgagtagcga gtgaacatta tctaaccncg acttatgtgc aaacgctatg ggatgactac 480
tccaaagat caccgagagag gg 502

<210> 1161
<211> 380
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 1161

atctaataatt catgtgagtt acagaaaaccc gcacttattt tgatcacact ctttgtata 60
cattattttt attattttt aaaacttata aaatcgagaa ttaatttcca ctgtatgata 120
agatgtgaag ccgacnattt tttagcctt cataaattgt aacagttcgt acaaatcaa 180
gcatattaaa atgtgttttgcatttca caatacataa tgaaagtggc ttcctgtcc 240
gtacaacaat ttgtcaacta gggcaagctc acattaaata tatggcatg gctaacatga 300

ttatcgaaa gtttctaaca ttatattac ccgtaaacta ttatgaatgc ataacagaaa 360
tatcaggaat attgttgtt 380

<210> 1162
<211> 424
<212> DNA
<213> Glycine max

<400> 1162

agcttaaaa gtacccattt accaacctca tagtctactt cacgccattt agtatcatca 60
gtgcgtttca tcgttagcttgc cgcttctaat agttctgac gaatgcaaag aaaaacttcc 120
tcacagtgc tcagagtatac atccaccgca tctagttgg aggagctagt gatatagtct 180
ggaaaagaga aaggtttccg cccaaacgtt atctcatacg gcgtagagcc cgtgcctgcg 240
ttccatgaag tattatgcga taattcgacc cagggaaagga atctgccccca agtccccgac 300
cggcggtgca ccatagcccg caaatattgc tctattactc tgttcatgac ttcactctga 360
ccatcactct gtggatggta ggctgaactc attctaagcc ttgtaccact caattaaaaa 420
agct 424

<210> 1163
<211> 427
<212> DNA
<213> Glycine max

<400> 1163

taggctgttt acaaaatcta tttagttaaa tatgctacga tttagtttat aataaaaacct 60
attaagcttg ataaattggc ctatatttc atttatatat aataaaaaat taatatacat 120
gtattatact ttaatattta atatcttaat aagttataaa ttcatcatcat aaataaaata 180
aatatttgag ataaaaagcc tttaaatggtaa taataggta tatcaggttt ttaaaaagg 240
caaaccaagc ttaaaaaaaag tctctgatag gataatagg taggtcaaac cttaaatttt 300
tataataggta caaacctatt tacacagagt ctaacatgcc ttgtatattc tcaccctat 360
tcttacgttt caatgtttga acttcaaaaaa gaaaaacaat attcataata tttgcttcca 420
atcatgg 427

<210> 1164

<211> 463
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 1164

 tcctcaacat caaaagtaata caacatccaa tcatcatgga ctatcaaaat caagcaaaac 60
 agggcaaagg cagaaaactc tgcccaaaac acaactcana atcacagctt ttctcactta 120
 aagaccagg taaaaattcc ttcggtccag tttgttaacc gttggatcga ctcgaaaatt 180
 ttactggaag tctctagtagtac ataaacctac attntgaccg ttggatcta ctagaaaata 240
 tccagaactc cttctgcaat attcttcca cagccaatca cacacaagca ttttctgca 300
 cttgtgc当地 attctgctgc acaatttcac agcaaaaatc tgcacaaaaga gcagatttcg 360
 aaaaccacac ttccccat ccaatcttc ccaaattcaga tcctacaagt cccaaatcat 420
 gtatcaatca tgtctaaacc aaagtcaagc ttcaaaacac agc 463

 <210> 1165
 <211> 401
 <212> DNA
 <213> Glycine max

 <400> 1165

 agcttggttc gaggtactca cccgttgaag atcgatgatc gactatgaac gaatgaagag 60
 cgtcgaataa cggttgaac ctttgcgaga ttccctcacgg aatacgttac ggaaacgttt 120
 cggaagcgcc tcggcttaga ttgtcttcac ggaaacaatt ttcccttagca tattcgaaag 180
 agagagaagt gcctattgtg ctgaacccct ttcttattgc ttccctcccc tatttatagc 240
 taaatagggg aggtggttgc cgcccagctc gcccaggcga gctcaactcg ccctggcgag 300
 cagggttgct tcctccagaa gctaccgcct tctggaggaa tattccagag ggcccaagtg 360
 ggcctgggtg ctatctgcac ccacattgta ctaagtacac c 401

 <210> 1166
 <211> 460
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 1166

cgctaaggcga gatctgcatt atttaaatac atttttcagg gtcaaaaaca caatttctc 60
atccccctct ctc当地actc caccaaaccac ccctagaaac tcctcccca ccacccagga 120
ccatcggtag ccaccacaag ccactgttgc ttgcccac accaccatac ggagagggaa 180
ttaaaatgca cagcggacat taaaagagta gggagaatg agacaaacac acaagagtnt 240
gtataactggc tcggtaacaa cccgtgccta catccagtcc ccaaggcaacc tgtggcctt 300
gagatttctt tcaaccttgtt aaaaatcctt ttacaagcaa agatccacaa gggatgtacc 360
ctccctgtt ctcttgaac ctatggatg taccctccac tagaactgat ccacaagaga 420
tgtaccctct cttgttctca gtcaataacc caagtagatg 460

<210> 1167
<211> 586
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1167

gcaaaaactcc caccatacgc ttctactaat catacatcac tctcaatatn acatanaccc 60
tcatacacac gccgtgcgtg aaaccattgt aganccgtcg tatctccaaa gacgacctcg 120
angcatgcaa gctgctactt atgcggcagg gccggacttac tttactatct tgtctccaaac 180
gcgagctatg accactgttc ttccttcccg cgacgcttct tttcatgtcc gcctgagtgg 240
gtatatagcc taaaccatac tttccacgat ctcccttgagt attcaacttag gctagttatg 300
ctgacgctgt ctgcctaa acccatcccg ggtgcataac cgctacacca acataactag 360
ggccattatt accgctgcat cggacagact aggctgctca aagagggagt ccacggagga 420
tatgtcgacc acctacaaag actgttagagc gtgctctaac gaatctatttgc cgatcacac 480
aaatgcaagg tagacggtca gcttaacaat atatgataca tgcccgatac gaagcaaaca 540
atgcctccccc actaccaaatttatttgc gcccagacgc aatccc 586

<210> 1168
<211> 380
<212> DNA
<213> Glycine max

<400> 1168

taaacagaca attttaaaga catgaatgta atgagatgg agtcttgatt tgccataattg 60

cattcaaaac tctcattgga aaagaaaatcc atgtctatga acttaggatc aataatggaa 120
tgagaggaga aaagggttgc gtaccatatac cgttgttctt ctgatgagaa catcaaggaa 180
gaagatatgg acgacggaat ctgcattcc tgagcctcg agtgcgcgtt ggcttcactc 240
gaagatccct tgcaactttt tcatggatct gccattgaa cgagttatgg gaaatatcaa 300
tcggctcacg tgaaagagaa tgacaacaga tgaagttgg gcttcggtg gagtgatttg 360
gacaacactc tactgatata 380

<210> 1169
<211> 427
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1169

tgttgtcat catcttaaag ggggagaatg tgaatgtatg tatacatgat tctgatgatg 60
tcaaagaaga atctaacaag gctgcttcaa atgataagca tttgcttcaa gaataattca 120
agattgcttc aacaaacaaa gccttggttc aagattcact aaagaccaag ccttgcctta 180
aaacaaagtg cttcaagac atgcaaggct ctggtaatcg attaccatga agtgaatcg 240
attaccagaa gacagggttg agaaatagct gttgaaaaag gtttgaatt tgaattttca 300
acatgtaatc gattaccata tgtctgtaat cgattaccag caacgaaact ttggaaattc 360
aaattcaaaa gtcataaccc ttcaaattat aactgtgtaa tctgatacac aaacattgta 420
atcgatn 427

<210> 1170
<211> 462
<212> DNA
<213> Glycine max

<400> 1170

agctcgtttat taaatacaaaa acacacatat tattatgaaa aaattgacgt taatgacgta 60
aattattatt aacacttacc actgcattgtc tcagctagtc gacatcagac cttgcagatg 120
tcgacgggtgc tgctgcctcc gtgaccggac ggatatctgt gtctggatcc tgaggggcaa 180
ctctgggctg ctagcatga ccatctgccc gaggatctga tggctggccc ggcgtcatga 240

atggatgcga catgcggaag aatcagtcca tgtaatcgct ggcacactgc cctggcacaa 300
cgcagatgtc acctgctaca accatatggt ccgaatagtg catccacctg ttgtgtata 360
catcagacgt gacccatgaa tcggcaggtg gagcatgaat ggtctgagtg tatcaaactg 420
ccgcatgacc ctctttggtc ggtaatataa aacaatgggc cc 462

<210> 1171
<211> 512
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1171

cgcgcgttaa tccgttggc annaccgtgc tcgtaccctg ggatactcta gagagatacc 60
tgcaaggcacg caagcatatg atactgagag atggaaaaca ctatttaat gacactaata 120
gggacatagt gttgtattgt attacaccac gtaatcccc tgagccttagc acactttctt 180
aacttatgct ctcgcgaatg aactataccg ttatttagtg agccttgaac agataaccac 240
tccctctagg aacaactttt tgctcatgat gcactatggc ctgtgtatag aagctatgtc 300
tcattcataa aatatgcttg aatcttgtat gctacgctac acctcactta cacacaatac 360
ttttgtaata ctttcattaa acgccaatcg tttcatttgt cggtagattt attctgctta 420
gcaccctaca tcccttacaa aaataggcac catttacttt tttgatccta cttcatcacc 480
tcgctctatg acggctatca tgaacaataa cg 512

<210> 1172
<211> 390
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1172

tgtcttctgt ttatatatga tttatactcg atctaagact tgtctgatgc aatttgctca 60
atcnntggatg acggcaatgg tgatttcgaa aatctgcact tatatgcaga attttgctgc 120
ttaagtgtgc atcgtaatct tgtgtttgtg cagaaaatgc ttatgcattt ctgtgtgtgg 180
aaagggttgt acatattggg gtctggacgt tncctaacat atcccaacgg tccaaatgt 240
gacttatgct ctatggacct ccactcaaattt tttcgagtct atcacacgat gaacgagttg 300

gaacgatcag aatgttactg gggctccga gtatgaaccg ctgcgggatc ttttgtt 360
ttggcaggg gtcttgctt ctgccctatc 390

<210> 1173
<211> 428
<212> DNA
<213> Glycine max

<400> 1173

ggaaataat aatctgccgc agcgatcca agagaaaatt aagagaagaa ggagaccctc 60
tgaaggctat ctatgctacg gtatgattaa ccttaccca tcgactttac ctttatgcc 120
aattgtaat cttaatcat gaaggtaag agaaggagcc tgtgttagtgt gtagaggagg 180
gaaaataagg gatgcctata aaaattatga atgaaagctt ataaaatatg aagtacaaat 240
acatacaatg ttgcacagat ctaaagataa tggcgaagag tgtcattagt tgatacttt 300
ttgttatact attctgtat tctgttgac tttatacgt ttctgtcagg cgtcataaaaa 360
aagaccgcac atagcttagga agactggtgg atgcgatagc ctatatacgca gtgataggca 420
taaacaca 428

<210> 1174
<211> 118
<212> DNA
<213> Glycine max

<400> 1174

ctataccac cgagcctga gatacgagt cattgcaaa catcatatac tacttgctgc 60
acaactatac ttgcttatga ttgccaacta gttattacat taatgtaaat tttctgtc 118

<210> 1175
<211> 483
<212> DNA
<213> Glycine max

<400> 1175

cgcgatgaac catggatacc gtgaataccg gaacctcgga gactcctgcg gcgtcgagtt 60
aaaaaaagacg agtttgcatt tttgacgggc gaacgcgtcg catgatcatg aaaccttagac 120
ctccttgcac taatgtataa ggtatgctaaa aagcataacta ttcgttccaa caatgactct 180

acactgctgt atctcacctc gccttggc atgatttact gactaccacg ccctgccttt 240
gaaaggatgg atttcaacac ttgtcacgct ctggtaatcc gttaccagga agagttatcg 300
atcaccagaa caccaggccc ttaattatca gctcaatcag ggtttgatt tttatctgca 360
acatgtcatc tgataaacacc tgactgtaat ttagtaccac tcctgagctt aggaagctac 420
gttctaaca caaaactcct caaattattc ggggttccg acacaccatc tctgataccg 480
gcg 483

<210> 1176
<211> 424
<212> DNA
<213> Glycine max

<400> 1176

ccaggcgacc tatgttgctt cccctagaac gcactgtctt ctggaggaac ttccctggaag 60
gcccaagtgg gcctgggtgc tatttcacc cccgtttac taaatacacc ccctgccttt 120
tttgctgat tcttttccg taacgttatg gaactttacg aatttcataa cgatacttgt 180
tttcttccg taatgtcacc aaaccttaca gattacgtaa tcattccctt ttggcttcc 240
gggatgatac gaaacttcac ggatttgca acaatgctt ctttgactt acggcatgtc 300
acggaacttc acggattgcc taacgatggg tgccaagtac ctcgaagtgg tctaaccgagg 360
gtcgtcatcc aacaaatata tggccccgg acgatatatg ggtatgacag ttgccccctct 420
ttat 424

<210> 1177
<211> 401
<212> DNA
<213> Glycine max

<400> 1177

tctacttatg tggcacggcg ggcttacttc actttattgt cttcaacgcg agctctgacc 60
actgttcttc cttgccgcga tgctacttt catgtgcggc tgagtggct tataatcctaa 120
accataactta ccacggtttgc tttgagcatg tatcaagcta attatgtccg cgctgtgtat 180
gtctacaccc atatcggtc ataaccgcgc cccaacatag ctcggccat cattaccgtt 240
gaatccgaca gacaaggctg cccatagagg gagtccacgg aagatatgct gaccacctca 300

taagactgga atgctgttct aacaattata tttgagcttc cacataacgc atggatgatg 360
ggaagcttac caagatgtct tcctctcctg acacgatgac c 401

<210> 1178
<211> 432
<212> DNA
<213> Glycine max

<400> 1178

agcttctgaa gaaggacttt actactctgg cgccagactag gagcgtcttg tcctacttca 60
accttgcct tacatcacat acatcttatac tgaacttaga tagggcgggg ttggtgtatg 120
gactagatat gaagatggat atgaatctt gaggccat ttctggacag atatctctga 180
tagctcagtc caactcctcc cggttaggat ttctagcgat tatcactgct ttatgcattgg 240
ccacaggagt caccttagac tcgttgactt tcaaaaactct cagcccagct attaacttgg 300
cttacatcaa gaagaacttc tagaacttgg atgacccttc ggtcagcttc ccagggacc 360
gtaaggccag ggccagagga tctgagggtc catcttcaac tgctccccag gactctacat 420
gtccagctcc cc 432

<210> 1179
<211> 427
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1179

tagtagcgt tacagaagtc aatatttccg ccaataacta tgatcatttc taccttacct 60
atttccatc atggccttgg tttgtgttt tagattgcca ccagagtcct aaatagacaa 120
tatcttcat tgcaagctta gcaacagtcc caaaacccaa ttttgccga aaccaagtgt 180
catgatttct atattacaa ttttgcttagc tggatgttt gcatcatagt tttgctatgt 240
catctacctt tggtctcatc tcttacctt acaattcagg caattatac attacccttt 300
ttcaatatat agaattggca acatgcaaac atatctaattc cagggaaattc caccactaat 360
agtcagccata taatccataa ccaatgaagt ccccatctc caatttattt catcatctaa 420
ttntattt 427

<210> 1180
<211> 407
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1180

ntcaacttcat aagttgtctt atgcctaana atacgttgag ttatcccttag taaaggtaa 60
aatattgact caagtaaaagt tcaattccca ctcaagattg gagcaaacta cttagcttgg 120
caataattcc gaccaaggc gAACCTAGTA ggtaaccatt cttgactgtg acgtatcaag 180
gaatgtattc ttaacgaata aagttacacg gttaagaatg ctgccgacag caacaacaaa 240
atggaaaatt ttgtttgtga cagaaaaatc ttttgtggac cacccatatg ttatttcaat 300
taatcttctt tttcatcgta aaatctttt ctgtgctgct ttcccatattt actctggaaa 360
atgaagaagc tatatacatt tagatgcagt tgtttattta tttattt 407

<210> 1181
<211> 345
<212> DNA
<213> Glycine max

<400> 1181

gcttcaacgg cgatttacca ccatggagat ggcgttggaa acaaggaga taaggtaaa 60
cgaggcgccc tctactacgg aataggccat ggaagacaga gcttcaccac caagaatgtg 120
ccttggataa gacagcttgg agaggatgct tccctggagg aaaagaggca cagattgaca 180
gagagagaga gagagaagat cgaccttgaa ggaggaacac gggagagaaa gttaaactct 240
gagttgtgtc tcacatgact ctcattcatc acatttacga caagcgatac atgtgctgct 300
atttatagac tacgcagcat acttgagaag cattctttag aaaaac 345

<210> 1182
<211> 421
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1182

tccatcgagt ggtaatcaga gcacaagagc ttcttagcaag tgctcccttac acctccatta 60
attttttttg ctttaccttc tcttccatttgg ttgtttcttc attttctcc atgtatctcc 120

| | |
|---|-----|
| tcacatgtct tgtgataaaat gttttaaca tgattctaga gtttccatgg attaaacttg | 180 |
| ctatataaagc tagagttcc aggatctta agctcgatg gaaagatctt ctggatcaca | 240 |
| gcactgcaat ttccttccat tatgatgctt tcctgattaa tataattgtt cttccttgg | 300 |
| aacatatctt tcaaaaattt atagtagtgc ggcattctgc acaacgcttc tctgaaggc | 360 |
| atggttatcc ccagttcctt gaaaatatct aacaatctcg ccagatgaca atcnntttct | 420 |
| t | 421 |

| | |
|-------|-------------|
| <210> | 1183 |
| <211> | 488 |
| <212> | DNA |
| <213> | Glycine max |

<223> unsure at all n locations
<400> 1183

| | |
|---|-----|
| gtgaacagtt tagttacaca attaattgta atctaannna tagccaggtg tgcatctctc | 60 |
| ccgcggatct taagtgaccg ggatgcactt tagcaagtgc cacttgtttt tttcttgaag | 120 |
| cccttaggc cggtgccctt ccttgtatta agccactaca gccctaaaag aaaacctgat | 180 |
| atcccgtatt tttaaggaat tggagcttag gaatagtca gaaaaatagag cggggaggtc | 240 |
| cttgccata gaaaaacttg taaagccgca atctttcag acgattttgg gccctcacat | 300 |
| gagtcctcg ttacgggaa aatgttggac atggcgactg gaagcgattc tctcaccg | 360 |
| ctgggtcgca accacgaact agagaagaca ttattacgcc tcacgtggct gaaggaatct | 420 |
| gcacggcca aatagagaga cttgatcca tcttcggaca aacctactga cctcatgacg | 480 |
| gtqqqatcq | 488 |

| | |
|-------|-------------|
| <210> | 1184 |
| <211> | 447 |
| <212> | DNA |
| <213> | Glycine max |

<223> unsure at all n locations
<400> 1184

ggtgtatngc tacattctac taatatatgg aattgcccac tgctatgcct gagaataaca 60
atngcttgac cacaacaacg ctggaggcg cg caagggacaa tggtc ttca aataaacctg 120
ttgtacatga acaa acatt a tatcatgcac tgaccgtgcc aaacgaacca gcgaagtcat 180

tgccataattg ttatactaac tatattcaat gtacctaacc aaaatgattt ccaaacacgt 240
gaccgacaca tatgtatgcgg tggccataag aatcaggtgg tgtgtgactt ctataaggaa 300
aaaatgtcat gtcttgttgg cgggacaacg atacaaggat tacgttatac cgaaaagcaa 360
tcacatatcc catgtccgtt atattcatcc actcgccac gcttacactga atgaaccaaa 420
catacacatg taagctaata taacatt 447

<210> 1185
<211> 388
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1185

agcttcttat caaggctcat cttgggtgg aagctccttc ttccatggct tattccttag 60
tggatggcgc ctcccttcac ctatttcct ttgtttccg ctgcacatctcc atggtgaaa 120
atcaccattg aaggacctca ttgaagctca aagatccagc ctccgtagaa gccccacaag 180
caagcttcca tcaagtggta atcagagcac aagagctca agtaggtgct ccttaaacct 240
ccatntaatt tttgctttac cttctcttcc attgttgntt cttcattgtt tctccatgt 300
tctcctcaca tgtcttgtgc taaatgtttt aacatgattc tatagagctc ccactgatta 360
aacttgctat acacgctaga tttgattt 388

<210> 1186
<211> 347
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1186

catagatgga tgaatgtatg ggacgaggtn gtacgcttagc tctagttcca taatggagac 60
cttctgctga aactaacttt ccacacttt gcacttcaca cgaaagagcc ccgcacgaaa 120
ctaggataga catgtccatg taaagacaga cgcttccaa ggaactacct tccgctcccc 180
atcatatatg gcaacgttct agattgctgt gcaccctctg atcttctagg ccattactgg 240
actgataatt ttgcgcgagc gacgacttca gctcatggac taccattata ctgattttca 300
tgagggaaata cggcgccctgc tgtggcatg actgtttact cccatgg 347

<210> 1187
<211> 376
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1187

taagtcacct gcggcatgca agcttctaaa ctttatacaa gaatgaagct ctgataccac 60
ttcttgaca agttgcctca gatatctaa gaagggggg ttgaattaag atatcacaga 120
ctattccccca attaaaaatt ctactttaa tttaatccaa caacccaaga ttcctttaa 180
acaagaactc ctagataata atgcaaatta atcttactaa atagaaataa taagcaataa 240
acaataaagg agtctaaggg aagagaaaat gcaaactcag atntatactg gttcggccac 300
acccttgtgc ctacgtccag tccccaaagca acccgctaga gaggttcact atcttgcaaa 360
atccctttac aagttc 376

<210> 1188
<211> 415
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1188

gacactctat agtactacag ctnttgttct gttacatctg tttcatgtat atgatgtgat 60
ctccaaatat ccaccgacta gtcatctta accagtgcaa ttttacatga tatcttgttc 120
ctgtagaatc tcatgtcctg ccgaagatgc ctgttcaaaa tttgccgtta cagtgagaaa 180
tccttaggaaa aatataagaa aacttgagga aaaagtgata ttatactctt cattaaaag 240
tagtacaaac tacaaaactta agtttagct tttagagcat attcattctc caataagtga 300
aaataatttgcgtcaggattt taggaagaat ttaattttct ttgatcgaaa tgcaaggat 360
tgctggattt cataacagta tttaaatttat gcaacagatc aaacatggc atatc 415

<210> 1189
<211> 412
<212> DNA
<213> Glycine max

<223> unsure at all n locations

<400> 1189

agctnnntgcg gattggtctn tgccagtgaa aggatcgatg tgggtccgaa aagaggcaaa 60
tntagtcatc ctgcttcgac gaatgagaaa actggngcaa atgaagaggg tgaggatgaa 120
agagaaaatcc atgctgcgt taccattcct acatggaaac ttcccaccaa cccaacaatg 180
tcattactca gccaataaca acccatctcc ttacccacca cccagttatc cacaaggcc 240
atccctaaat caaaccacaa aacccaccta ccacacaacc aatgctaaac accactttg 300
gcatgaaccg aagcaccaac caaaaggat tttgcagca taaagcctgt aggatccacc 360
ccaaattccg gtgtcatatg ctgaacttgc tctcatatct actcgataat tc 412

<210> 1190

<211> 470
<212> DNA
<213> Glycine max

<400> 1190

tcaagccagat cgctaagtga gagcttatcc gtggctaagc atgacctatt gtcgccaagc 60
gcaattcctt acgaccataa ttgaggtcca tgacgctaag caccagtcat ggcagctatg 120
cgagattcat tgtggcaata tgagcgctaa gcgagtcct ctcagctaag cgcatactcc 180
tctgtactta agatgcatca ttttagctaa gctggccaga gcctgttttgcgagagttg 240
tagctttct aatctacaga cctcgctaag cggacatacc ctcgtgctaa gtcgagttc 300
tgctaaaaaaaaaaa aaaactgatt ttgaatgtga aacgtcagct aagcgcacgg gtccgctaag 360
cgagccttgt tgagaaacca aacgtctctc ttgctcgctt agcacaacgg tccgctaggc 420
gaaagtatcg aaaaactgtc taagtgagtg taacagcagc tacactcaca 470

<210> 1191

<211> 474
<212> DNA
<213> Glycine max

<400> 1191

cggttgtgca tctactcgac cggatcttaa gtcactgggc tgcagctcta gccaatggac 60
taccttgatt aattcctttg gagccctttt gagccgtggg tgccttcct tggttgaag 120
ctcaactacaa gccctaaatg aaaaaccatg atatcaccct atcttgacg aattttggag 180

ctctggatt gttctgcgaa taaacgcggg gggcttttgc gtttcattgg agaccgtat 240
cgcttagcatt gttcgttat gtatTTGGG ccccacttga ttgacattgt atattggtaa 300
acgttggaca tgctgaatga aactgtgtgt ctcacaggct atggagtggg gcacctgctt 360
taagaagagg atttgagcg ccctaagttg agtgaataaa atcttcaccg tccagaatgt 420
gaaaccctgg tctcccttat gtaaaatgaa cgtaactttt tagttcttaa tttt 474

<210> 1192
<211> 456
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1192

tgcattgattt acattctccc tctttctcaa gagaattctt aattcttctt gacatcatca 60
aaatcttcat gatttacatt ctccccctt ctaagaaaa ttcttaattc tgcttgacat 120
catcaaaatc ttcatgattt acagcctcag ctggccttgc cttggctt ctggcccccac 180
aaataataat tggtggatca tcagggttcc aacaattctt cttatgtan gccagattaa 240
tgaccggct cagacgttca aagacaagac tgtcggaggc aactcctctg gtcctgcata 300
aagctgttat aagagcagga aacacgagtt ttgaggagtt atactggcc atagaagtaa 360
tctgccctaa aatgagggttcc ccgacattca tgtccagctg agtaactagg ccaaagatta 420
acctagctcg atccacagtc agatcagatg tgtggg 456

<210> 1193
<211> 504
<212> DNA
<213> Glycine max

<400> 1193

gcaccggatt gatgcgatcc tgtctccgag catgctatca ctctaaactgc acgcgtgcgc 60
atccccatatc gacggtcact ttcgggttat tgattttctt agcatggctt atgacatagt 120
tggaaaggccc ctactctcac ctataactcac tagacttacg ctgcatttcc gtggaggacc 180
ataaccattt gaggaccttca ttggttctca tatatcccc ctccgttagaa tctccacaaa 240
ccatcttcca tcaagaggat ctcaaattcac acgagcttga agataactgct acgttaacgca 300
gctcaatttgc tcgcttaccc tatcttcattt agcgctgctt attggatcta cgtgtatctg 360

ctcacatgtc ttgagactaa tgtcctaaac atgatattta gagcgaccac tgattataat 420
tgctatgtaa gcataatgtg atcgttatag cgcacatctc ttttcttgta tctaacatga 480
attggtagag taaaggcctt tagc 504

<210> 1194
<211> 417
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1194

agctntatgg ccatgtaaac actatggctt atggttgtt ttcccccatt caatcaaccc 60
aatgtttcca aaaaacatct cttttatcaa gtcacgcaca catccgagtc cattaggca 120
tccggaaaaa atcttcatt gcattcaccc ttcaggcgca cacattttt tcaaaaatct 180
ttttatatcc taatctgtga atttccaaa gaaaactggc ggtcatttc tttcaaaagc 240
atgttggctt ttttagtttc tttctcttag ctnnnnnn caattaattt ctttcagacc 300
aatttttttc agaaaagggtt tgtaacctgg gcaaagttgg tattcgagat tacactntat 360
caaaaggaac aanaggcgtg tgaatgacaa taaaccaaca cacaagaccc ctcttat 417

<210> 1195
<211> 426
<212> DNA
<213> Glycine max
<400> 1195

tttcgaagg gcaaggttat ttccagtttc ttgaaaatat ctaaaaatct tgccagatga 60
cgatctttt ctttcttggaa aggtaccaca ggatatagtt cttccacacc ttcatccaca 120
gcttttcac ttctactctt ctttgcattt ttatttatt tattttttt catttctat 180
ttcttattct acttctttt cttttcttg gtccttcaat tctttattct ggaccattat 240
ttgtttccct ttttcccgat tgccttcacc ttcacatca tttttcttaa cttcagtacc 300
tttctttta gtcgctttct ctttgtgcac tacacttct tcatttcag cttccacaaa 360
cctcttactc cttgccatca cagcttgca ttcccttgc ggattctgtt ctgtattgc 420
cacaaa 426

<210> 1196
 <211> 464
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 1196

```

tctccgcattttctataat ttagggggagg aagtgacagg aangaacgtt caaccctcct 60
ggtatatgag attcaacttac tatttagtgag ataaaatcgct tccatgaaga atatacacgc 120
cgaggcgctt ccgtaacgtt gatgttacgt tttcgtgagt gatttcgcga agattttcaa 180
ccattttcg acgttcttct tttgatattc gtcgttcttc ggtttcaac cgtaaagttc 240
ccgatatcga actttntaat tcattctatg taccgttggt ggtccccatt cgtttagcgt 300
acttttattt tcgttcata tactctacgt agctcctttt gacgtgctt agtcatctac 360
ttgcctaatac aataataaaaa taaattcca ccgatcattt gaatggttac atcacttaat 420
ttcagttcaa tgagatgtga ccgtttggc atgccataac catg 464

```

<210> 1197
 <211> 100
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 1197

```

acatgttgct ccccctatct ctaacaagct tctatagttc ctgtgatcta tatggatact 60
gngatagcta ctatatcgc cttgcgttga ggacgggcta 100

```

<210> 1198
 <211> 281
 <212> DNA
 <213> Glycine max

 <400> 1198

```

ttaacatcct cattgattag tgcagatcaa actactatat ctttgagggg acttacatca 60
tacacttgag ctgcattttc cataaaccag aattcgcaaa ggatgaaatc aaggaatagc 120
atacataaaag atcatgagca tacccacgtt ttcaaatttg accataaagg gccataaaaa 180
taattcttat atttcttaa aagatataattt atttataat aacacatagt tggtatagaa 240

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tgttgatgct tacaagagtt aaaaaattat tcttatccaaat t

281

<210> 1199

<211> 197

<212> DNA

<213> Glycine max

<400> 1199

taagtaaacg atcataaaacc cataaatctgg cgacaagtgc agataatgag agtcatggct 60

agttggcata acatgttaac caatgcacatct agtttacctt cacgcttcctt atttctgttt 120

atgaatatga attcacggct acttgattca ctccctctaattt gaccatagca tcacttctgg 180

cactaaatttggag 197

<210> 1200

<211> 413

<212> DNA

<213> Glycine max

<400> 1200

cacggagact aatcagacat gggatgcagc tatcacgtac atgcttctat tctaaaactt 60

ctcatcatgt gcttatttagc tggtcggtt tctcttggc tattgaagcc ataccaatta 120

tggacaatat tatggtaaca ggcagaagat ttccggatg attcatttctt gggatactat 180

ggtaagaact tggaaatctca tcttcaatga gtcataattgt cataccatga aatatcatat 240

cttggcata ttgaatgaat cttgttaatta ctacaaaaca cctactgaaa ttctaaattt 300

tttagatcaa atgctaagta caagtaaaaa gatggatgat gatactattc tgcaggcttt 360

agctgtgctg tgtacgcgtt tcctataattt gctattgcta tcattggact tct 413

<210> 1201

<211> 596

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 1201

cttcacactc ttntcaacat ttaantgtgt tgtnntnttaattnnt ttttnaggcc 60

ggaatgatgc ctcacactg acannctaaa gttaaccgca gcatgcgact taacaaccag 120

tagatatgag tgaattaaac tacgattaca agaccttcaa caaatgtgga agaatatgat 180

tgaagcatac ttttatattc tagcatatct acacagtaca ataacgtaca ctttggatt 240
aaggccatgc ct当地atctt tggctgctgc attattcgtaaaacatagc gcatatgcc 300
tataactaacc cggattact gtaagtgaac tcggaggcat tacaagaat gagaatttag 360
acagttctta gtttcgagtt gataagataa gtgcaccatt acaagggtgcc aacatgggt 420
cacacccaaac cgaaaacaat tacagcaata aaaacaaagg tcatgtacac atggatgtg 480
attatagctt tgtacatgga tggttctaag ataaaaagt acgcttgcca ttttacgta 540
atgtggtatac aactttggtc ttaggcccac cattacaaa cactacaatt gtcatt 596

<210> 1202

<211> 484

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 1202

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catctttat ttatTTaaa cgaacaatag taataattac tgtgaattta aaggatactg 120
ggctgcttta tagcagcgcc ttccgcttgc cccggctag gcaaaggacg acgacact 180
ggcgtgacc ctagcctctc tttgcattcg tccataagta cctaaaagta gaaaacaatg 240
agggtggca aatcgcgacc gcgtcgctgt cttaccttaa tcggttctg cttaactt 300
tgtctcaacc tttgatgatt ctgccccctg tttatcacaa aatatgcattt tgtatgcgt 360
tgcattaatg tttcaaaacg caacaaattt ttagtgaag ctggtttagg ttcatnta 420
atataaggcgct tggggcatcc catgaactga gcaaaaggc tcaggtgatc acaaactaac 480
acat 484

<210> 1203

<211> 441

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 1203

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ctatgctaga ttgcacacca tgtaacctga caacctcact tatatacgag gaggtcaact 120

tttccatgga aaatctgata ttaatggaa tgaattgagc aaacttattc aatctatcaa 180
caataaccca tatagaatct aaacctctaa gggttctatg tagtcctacc acaaaattca 240
tggaaatgct gtcccacttc cactatggta tctctaaagg ttgcaactta cctgaaggtc 300
tctgatgtta tatcttatcc ttctgacaga ctangcatgc atacacaaac tcactaacct 360
ctctcttat gtatagccac caaaacatcg tctttaaatc atgatacatc tgtggagcac 420
catcatcaat gctcaaatta c 441

<210> 1204
<211> 313
<212> DNA
<213> Glycine max

<400> 1204

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catgcgtctt acaatgtatt gagaatgtta ccgcctgccc ttactagctt ctgatgcgca 120
caagagagct cgaggatcat ctgcttgtat acaagggta tgctgtaatg catatcagat 180
agtctgttga gcgcgtatgc acgacgatgc catgactgct gcgcacactag atgctggtgt 240
ttgataacag acatgagcac gaatgatagg ataaacgtga tgtgattaac agatgcttat 300
gcactgcatg ata 313

<210> 1205
<211> 239
<212> DNA
<213> Glycine max

<400> 1205

agttaactga ggcgtgcgca cgctgtctt gaacttagt tcactctact gctgagcttg 60
cgccatcat ggaatatgca gatctaagca gaagctcgaa gtgattcatg atacagacca 120
tagctctact aagacgaaat aaataaatga gtatcaatca ctccaatgag catgctctat 180
ggtgacatca tggattatca cttggcgtga tgaagtgatc tgctcgctta tatattaat 239

<210> 1206
<211> 357
<212> DNA
<213> Glycine max

<400> 1206

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tctgtacaga ctagagtcgg aagcgaaagc cggtctgaga gctggagatg aatccgtac 120
tggtttgag atcctacagg tgaaggagac atcctcagca cttgtatttc tgcatctat 180
cgtcttgttc atagcttcg tgtaaaggac gtccttgga tatggaaagc tcgaatacta 240
tcatggatct tcgttgttg tgcacgatgt ctatataat ctatgtattt aatgaagtta 300
cgtgagatgc ctgtgctata aacattttaa ttcatgtatgc ctataccatg ttcatat 357

<210> 1207

<211> 458
<212> DNA
<213> Glycine max

<400> 1207

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gagtttgct gaaagatatg tactccttct gcctatgtaa attatgttt atgtgattat 120
ttataactta tattggaat tgcgatatat tgcgtgatgtt ataaggctga tattattgg 180
atcattattt tttttttttt gaaaatttac tcttttcta tctcgtgttt aggagaattt 240
cttatactat ttacttctct ttagtagtac ctattgcttc cttgttttt ttttttttac 300
ctcttgaact cctccgattt cgggtctctg aagataatat catgcgtgc tggtaaca 360
atatacatgt ctgtgctata gtcattattt acggagat gtatggacta caatataatg 420
gctaagcgta agaagaggtt aacactatta tatataat 458

<210> 1208

<211> 517
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1208

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ttgatagtaa aaggctgaga tgaatgtca accttgatat gcagaccatg atgaagagag 120
cctntgaggt accttacaat gcgtataaca gctgtccaaat gagagtccat atgattagcc 180

atgatctgac agaccttggtt cacagcatag ctttaactcat gccttgtaat ggctgtgtat 240
tggagagcac caacaacata cctatagaga gatggatcac tgaataaaatc caagccagct 300
ttggttaact tgcaattagt agtcataaggga gaaggaatga ctgtgcttct gccatTTGG 360
ttttctgaag taaatctctg atatagttgc tgagtcagta gaatagtccc atcagccaca 420
gattggattt ctataccaag aaaatattca áagtttccaa ttgtgttaaga caacaattgg 480
aatgtatctt ggtggtgagt tgctgaatta ttacatg 517

<210> 1209
<211> 366
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1209

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atgtcaatac caaccacaga gagtccaatt aagacatcca tagttcctt agtttcacaa 120
tatcatatct taagttggtt atatataaat atcactatca tagcttaccc acacagtgac 180
aaagcataga catacactga ctctgtcaat ttattccac tatgatggga taaaaaaactt 240
ggcattatgt tcagttctac tatgtctcta ttgttcttctt ctttctatgt gcttgttaat 300
caatTTTGc atgtactttc aactgtataa atangaaaat aattttcctt tatgggtctc 360
ttcttag 366

<210> 1210
<211> 254
<212> DNA
<213> Glycine max

<400> 1210

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tcatgagtagtac ttcacacaaa actagagggg ctaccacagg ttgcttcaac accatgtatt 120
ttgctgttat ctccaccact tatgatccta gtgatgaact ctggcatgac cttaatagca 180
acccttccat cacaaccctt catgtactga atagggcgtg caccagactt gatacgggac 240
ttgttagggc aaga 254